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qMRI analysis of cartilage and bone.

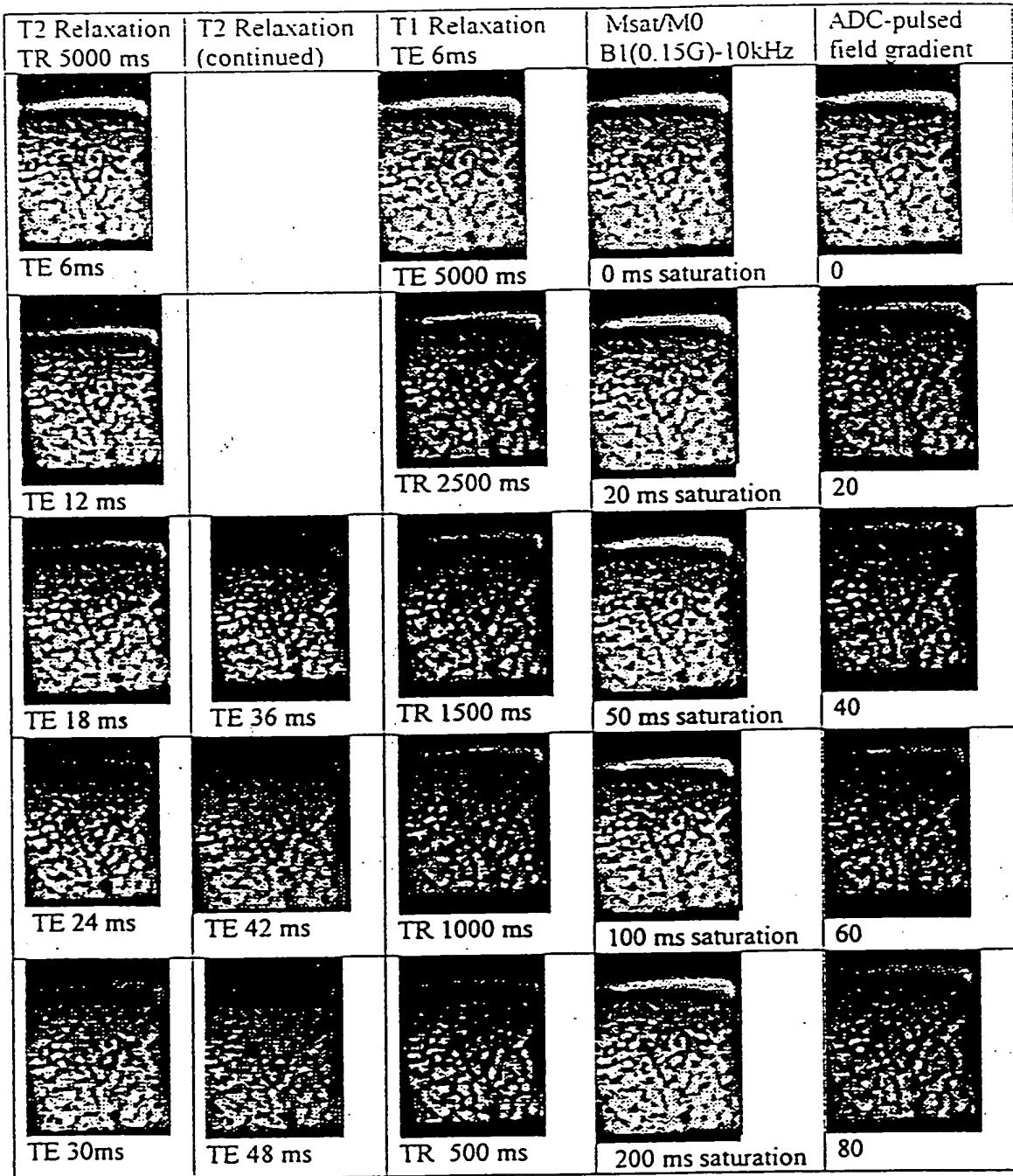
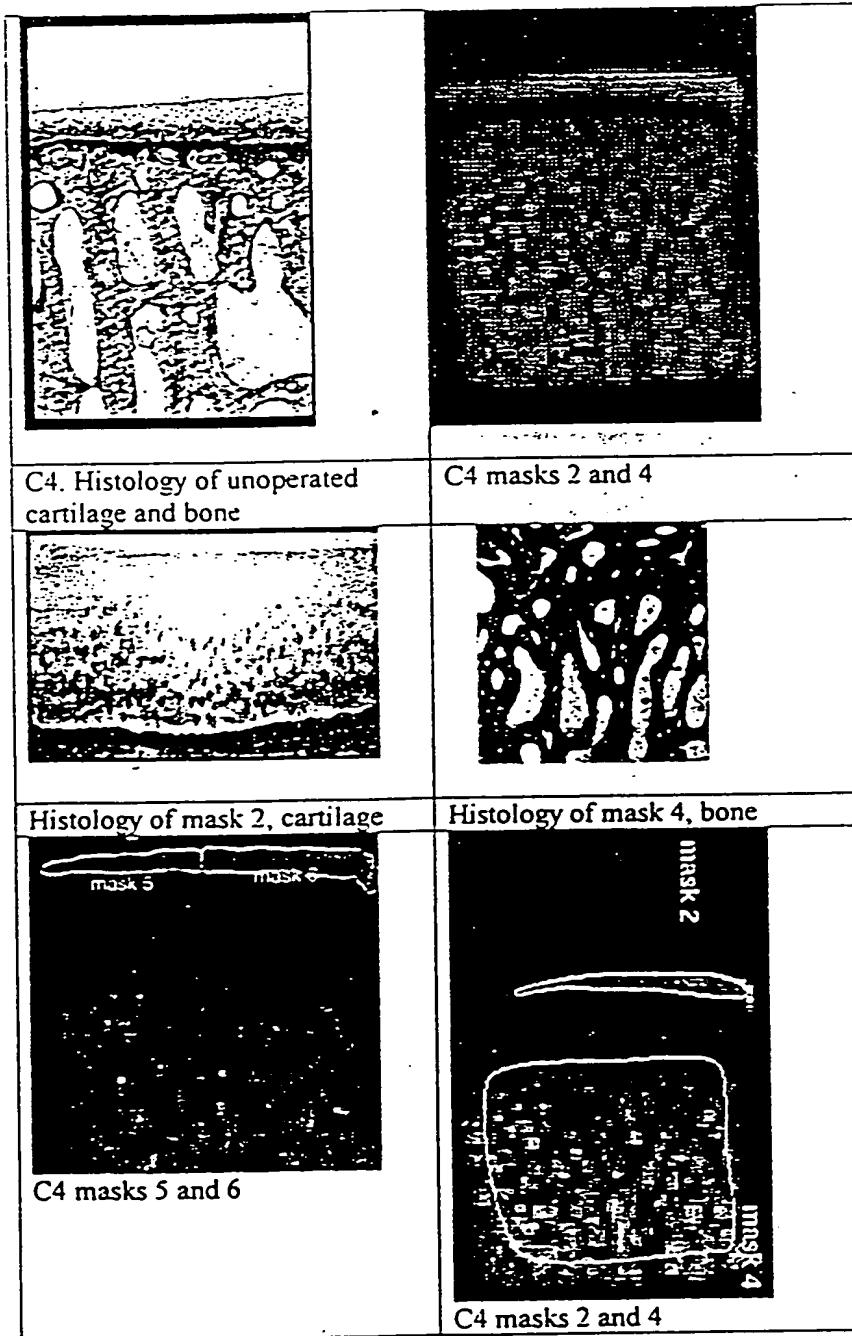


Figure SRL 1

A complete set of MRI scans from the qMRI analysis of one osteochondral plug (C4) showing the sequence of data acquisition and change in signal intensity of the cartilage.

09828070-04150E

C4 Unoperated control



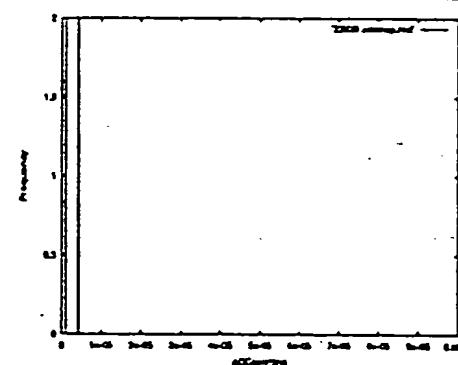
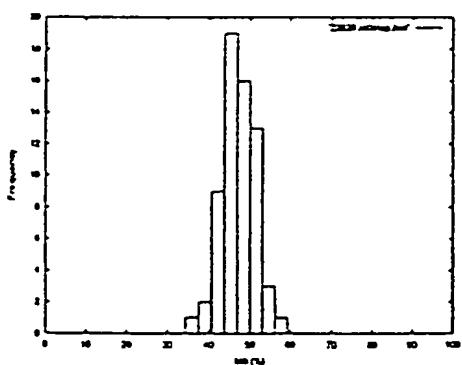
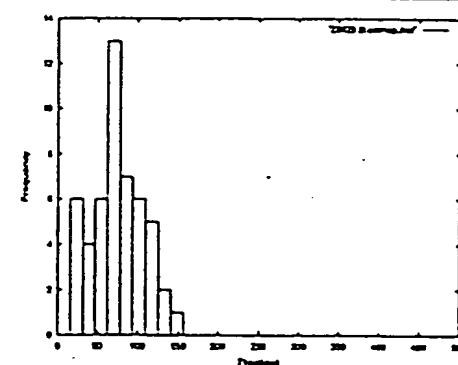
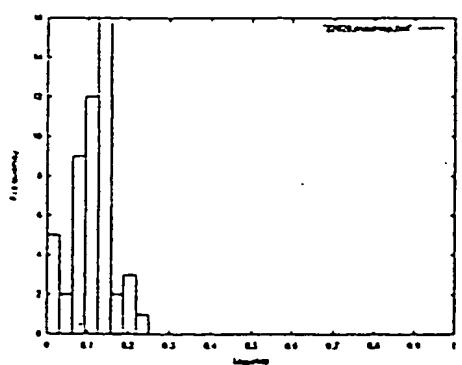
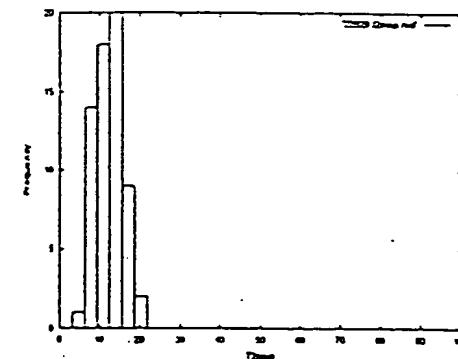
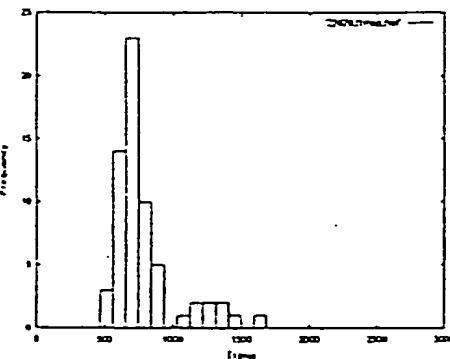
00000000000000000000000000000000

Figure SRL 2

Definition of masks 2, 4, 5, and 6 chosen for qMRI analysis of cartilage and bone.

## C4 mask 2, unoperated cartilage

Parameter	Value	(s.e.)	Parameter	Value	(s.e.)
M0(%)	48.8	0.5	Msat/M0	0.116	0.007
T1(ms)	812	30	T1sat(ms)	77	5
T2(ms)	12.8	0.5	ADC(cm <sup>2</sup> /s)	8.1e-07	3.1e-07



Picc 3838-3843

x axis - number of pixels with the same value

LHS- top T1,middleMsat/M0,bottom M0.

y axis - range of values.

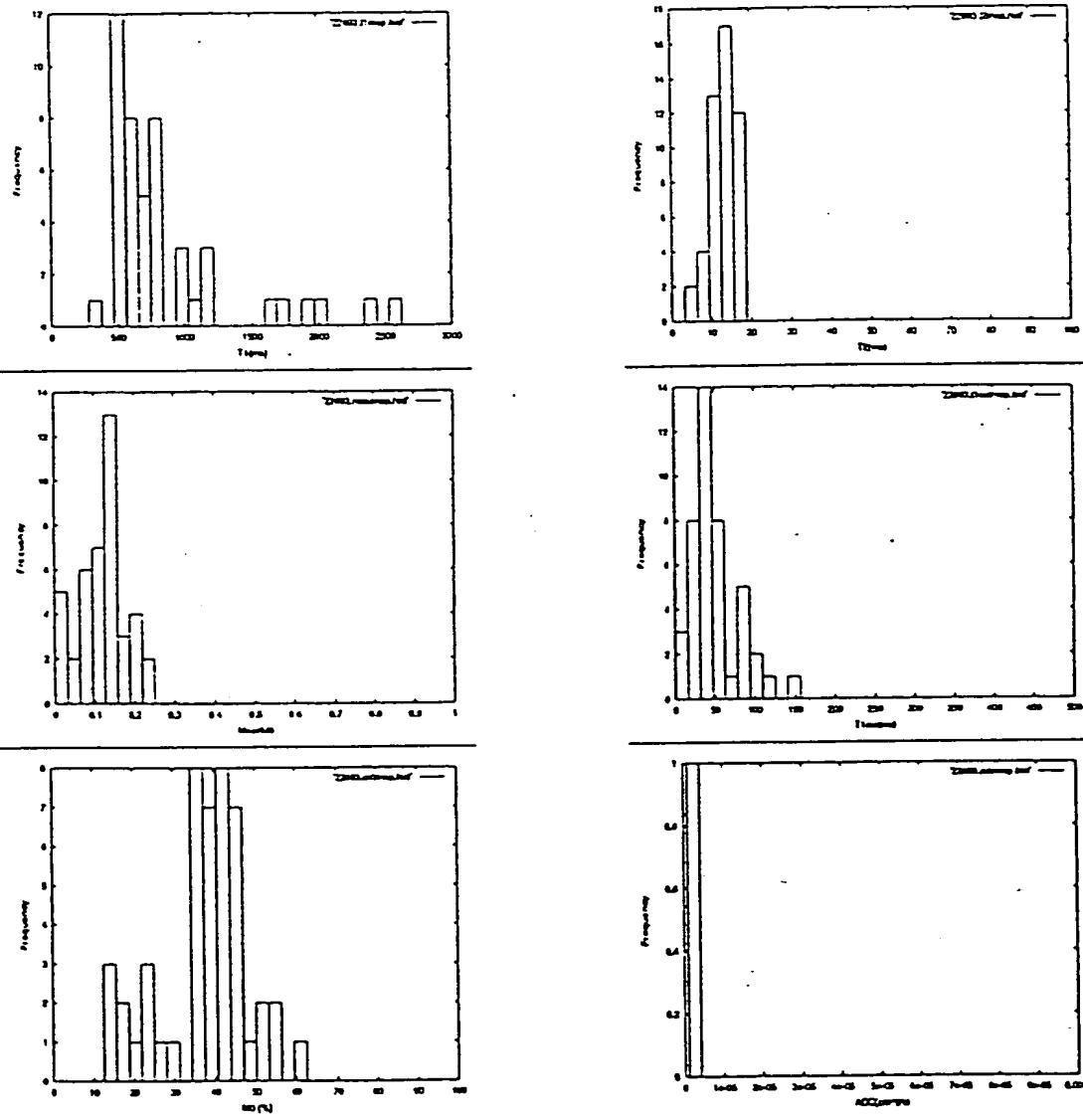
RHS - top T2,mid T1sat,bottom ADC

Figure SRL 3. Automated output of histograms for C4 mask 2

105410-02082860

## C4 mask 5, unoperated cartilage

Parameter	Value	(s.e.)	Parameter	Value	(s.e.)
M0(%)	38.5	1.7	Msat/M0	0.125	0.009
T1(ms)	907	76	T1sat(ms)	53	5
T2(ms)	13.8	0.5	ADC(cm <sup>2</sup> /s)	6.4e-07	2.1e-07



Picg 3838-3843

x axis - number of pixels with the same value  
 LHS- top T1, middle Msat/M0, bottom M0.

y axis - range of values.  
 RHS - top T2, mid T1sat, bottom ADC

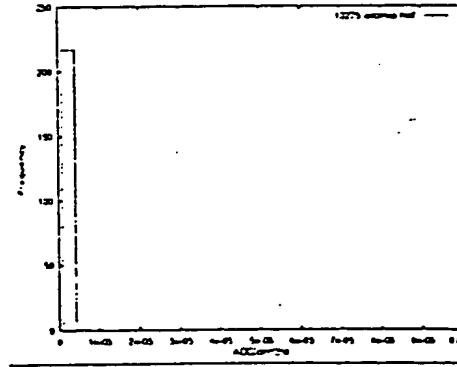
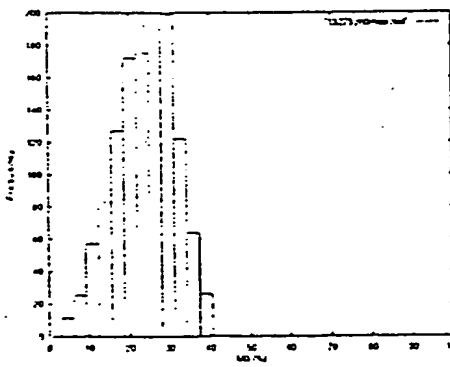
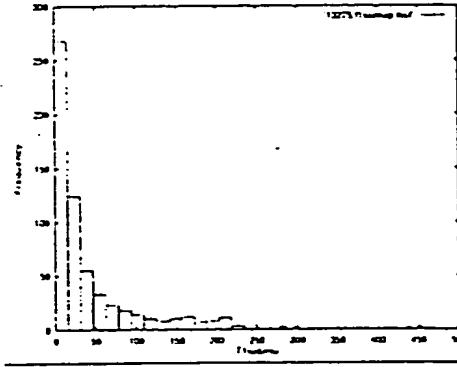
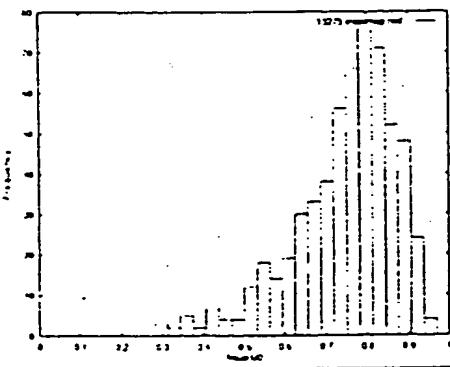
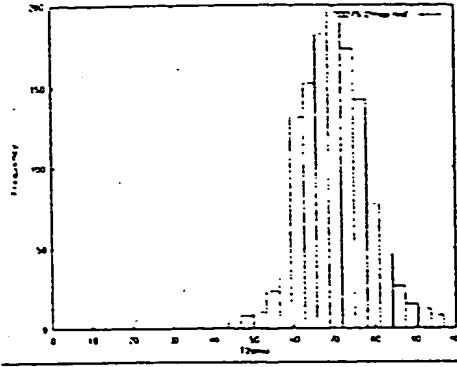
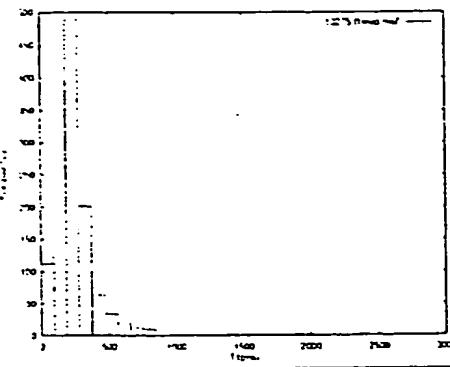
Figure SRL 4 Automated output of histograms for C4 mask 5

105010-020828160

## C4 mask 4, unoperated bone

Parameter	Value	(s.e.)
M0(%)	24.9	0.2
T1(ms)	291	9
T2(ms)	73.2	0.3

Parameter	Value	(s.e.)
Msat/M0	0.769	0.006
T1sat(ms)	63	4
ADC(cm <sup>2</sup> /s)	8.6e-07	2.4e-08



Picc 3838-3843

x axis - number of pixels with the same value

y axis - range of values.

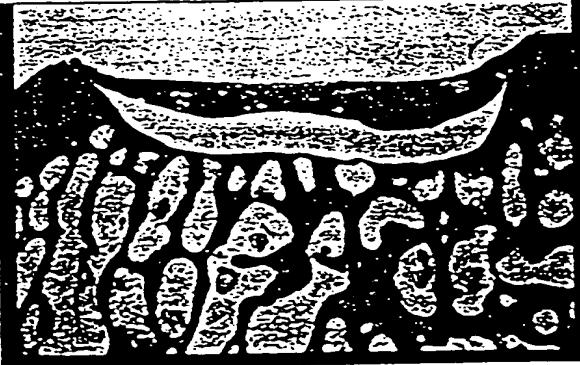
LHS- top T1, middle Msat/M0, bottom M0.

RHS - top T2, mid T1sat, bottom ADC

Figure SRL 5 Automated output of histograms for C4 mask 4.

105010-0202860

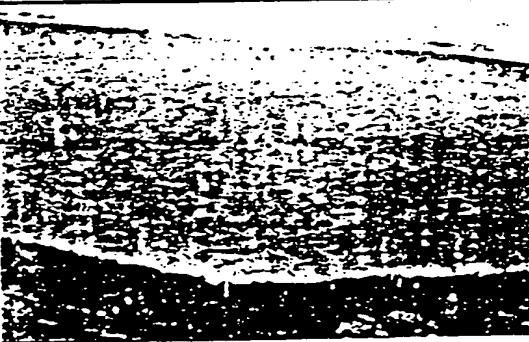
C128 (ACDI + HAED, vital flap) zero time point



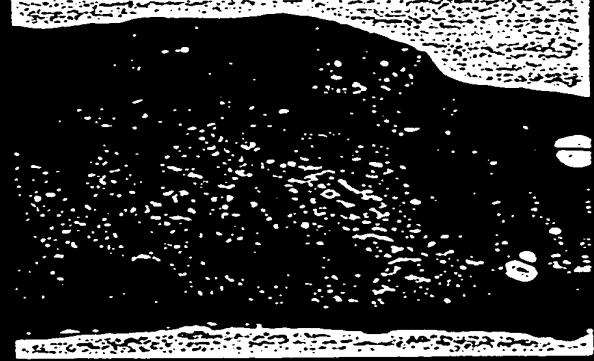
## C128 Histology of defect



## C128 Histology of defect



### Histology detail of mask 2, operated cartilage



## Histology detail of mask 6, synovial flap



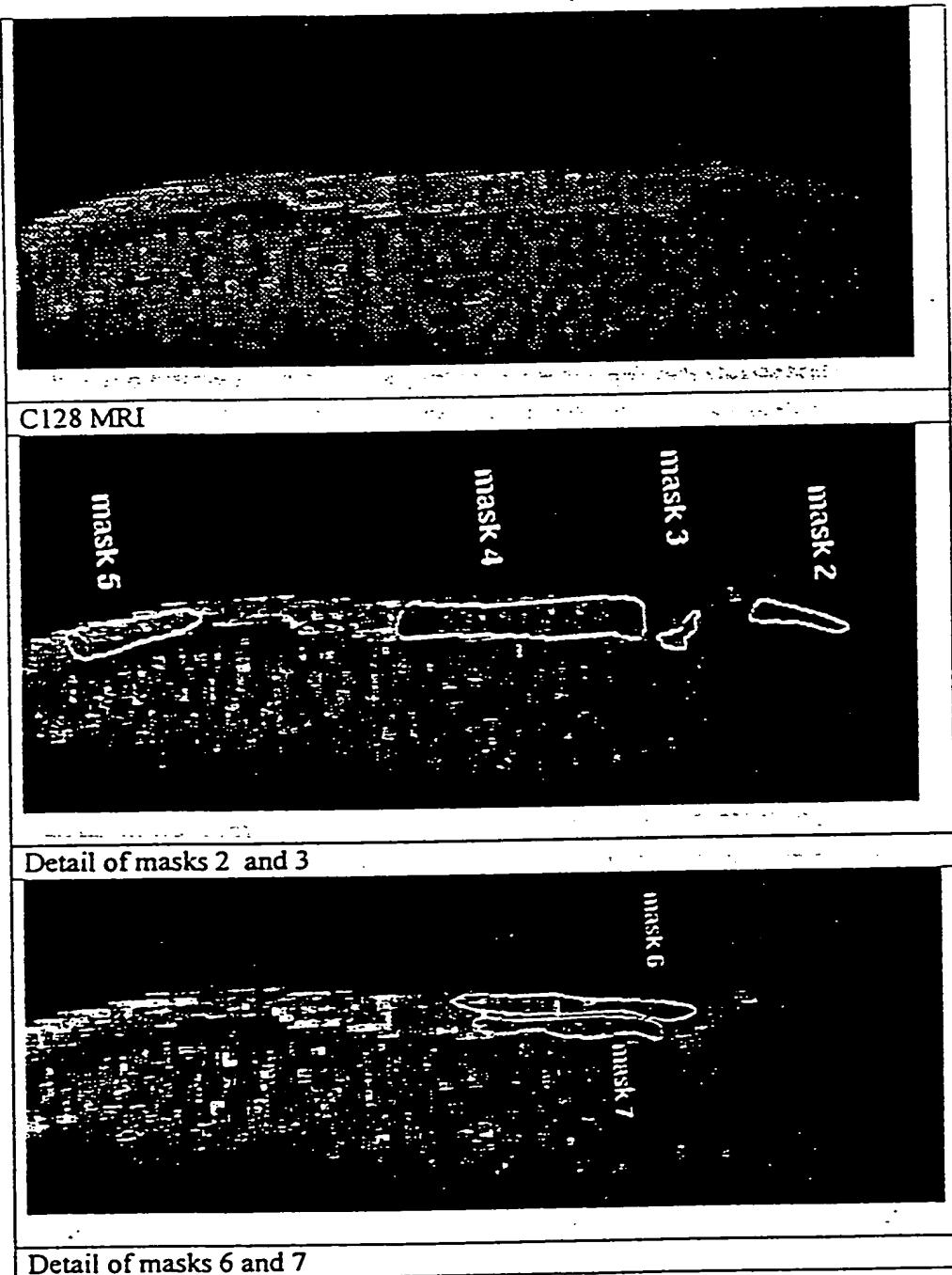
Not available – presumed lost during histology processing

### Histology detail of mask 4, HAED matrix

### Histology detail of mask 7, HAED + synovial mince

Figure SRL 6. C128 (ACD1 + HAED, vital flap) zero time point  
Histology showing detail of masks 2, 3, 6 and 7.

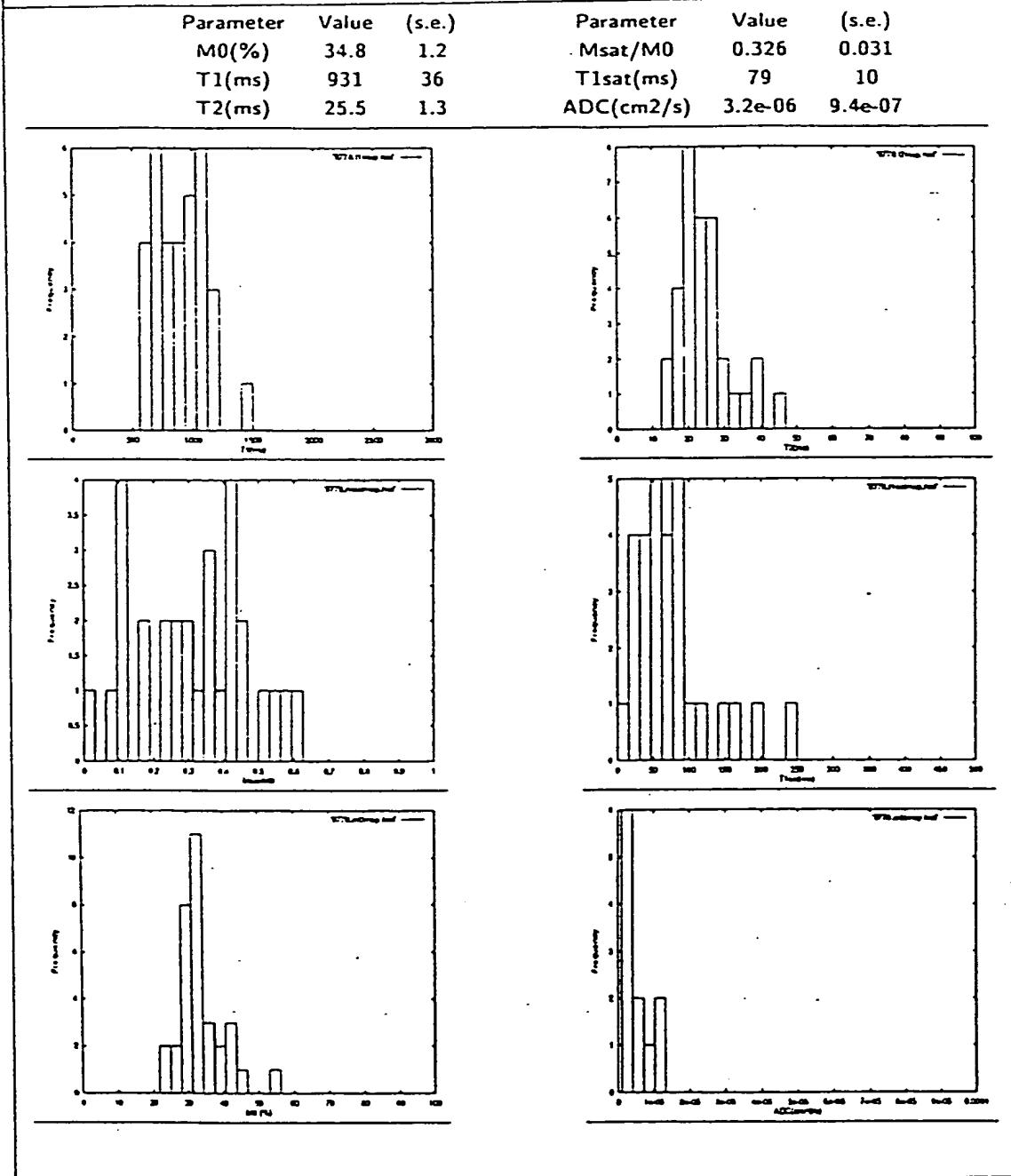
qMRI analysis of C128



Detail of masks 2 and 3

Figure SRL 7. C128 (ACDI + HAED, vital flap) zero time point  
Definition of masks 2, 3, 6 and 7 for qMRI analysis.

## C128 qMRI values for mask 2, cartilage adjacent to the defect



ongg 493-498-2

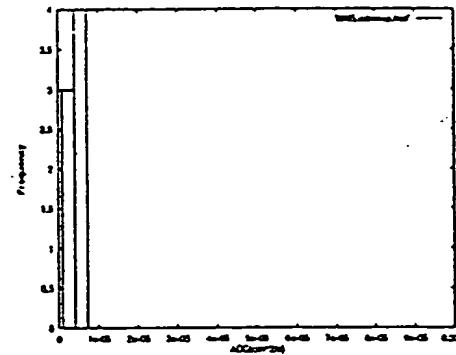
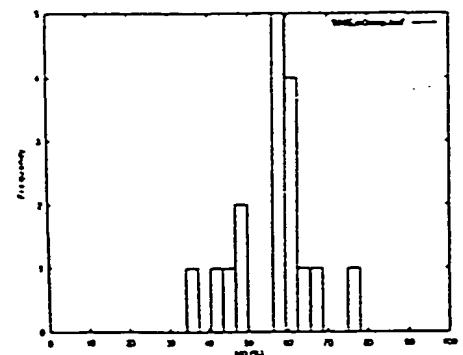
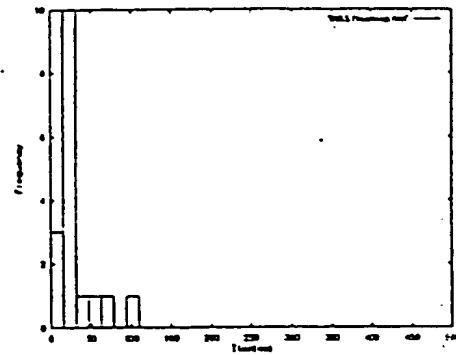
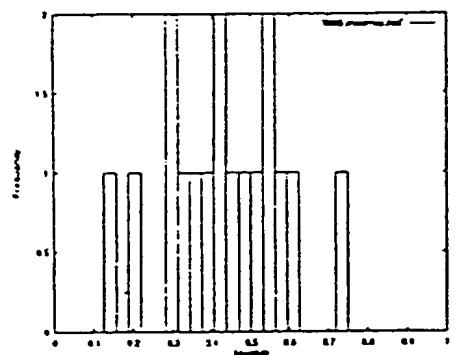
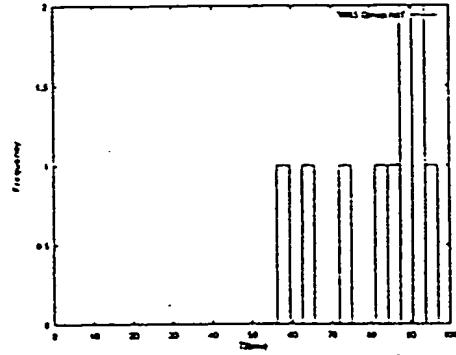
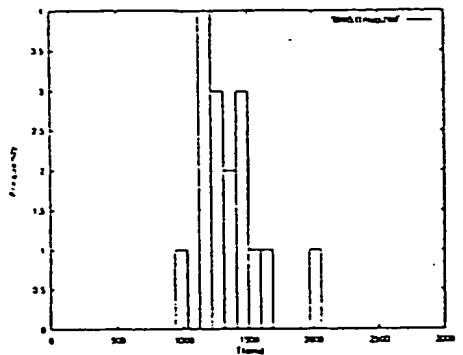
x axis - number of pixels with the same value  
LHS- top T1, middle Msat/M0, bottom M0.y axis - range of values.  
RHS top T2, mid T1sat, bottom ADC

Figure SRL 8. Automated output of histograms for C128 mask 2

## C128 qMRI values for mask 3, HAED matrix

Parameter	Value	(s.e.)
M0(%)	58.5	2.5
T1(ms)	1684	200
T2(ms)	96.9	4.2

Parameter	Value	(s.e.)
Msat/M0	0.442	0.037
T1sat(ms)	31	6
ADC(cm <sup>2</sup> /s)	3.4e-06	6.3e-07



long 493-498-3

x axis - number of pixels with the same value  
LHS- top T1, middle Msat/M0, bottom M0.

y axis - range of values.

RHS - top T2, mid T1sat, bottom ADC

Figure SRL 9. Automated output of histograms for C128 mask 3

## C128 qMRI values for mask 6, synovial flap

## Parameter      Value      (s.e.)

M0(%)      48.7      1.4

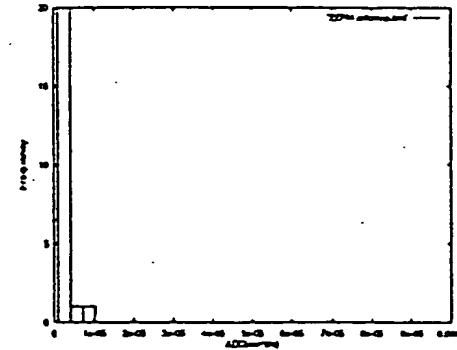
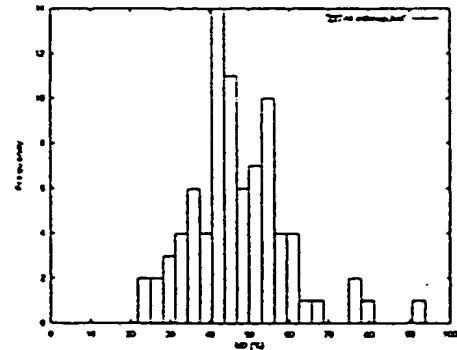
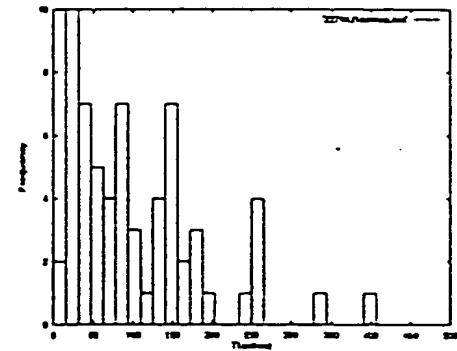
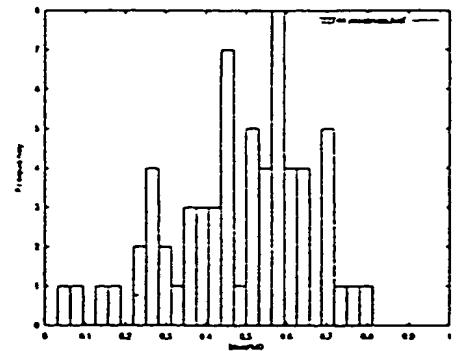
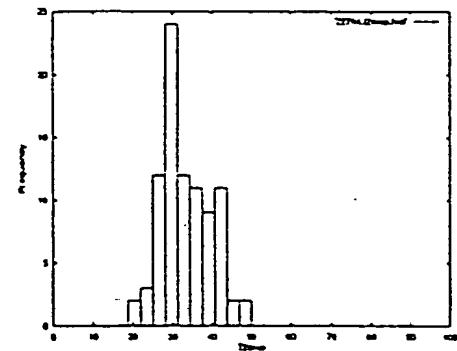
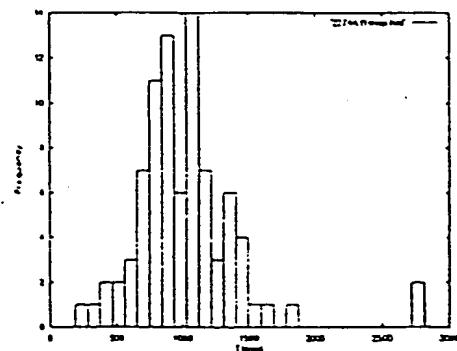
T1(ms)      1083      52

T2(ms)      34.5      0.7

## Parameter      Value      (s.e.)

Msat/M0      0.496      0.022

T1sat(ms)      110      11

ADC(cm<sup>2</sup>/s)      2e-06      3.2e-07

ongg 493-498-6

x axis - number of pixels with the same value

LHS- top T1, middle Msat/M0, bottom M0.

y axis - range of values.

RHS - top T2, mid T1sat, bottom ADC

Figure SRL 10. Automated output of histograms for C128 mask 6

F05000 "OCT0820060

## C128 qMRI values for mask 7,HAED + synovial mince

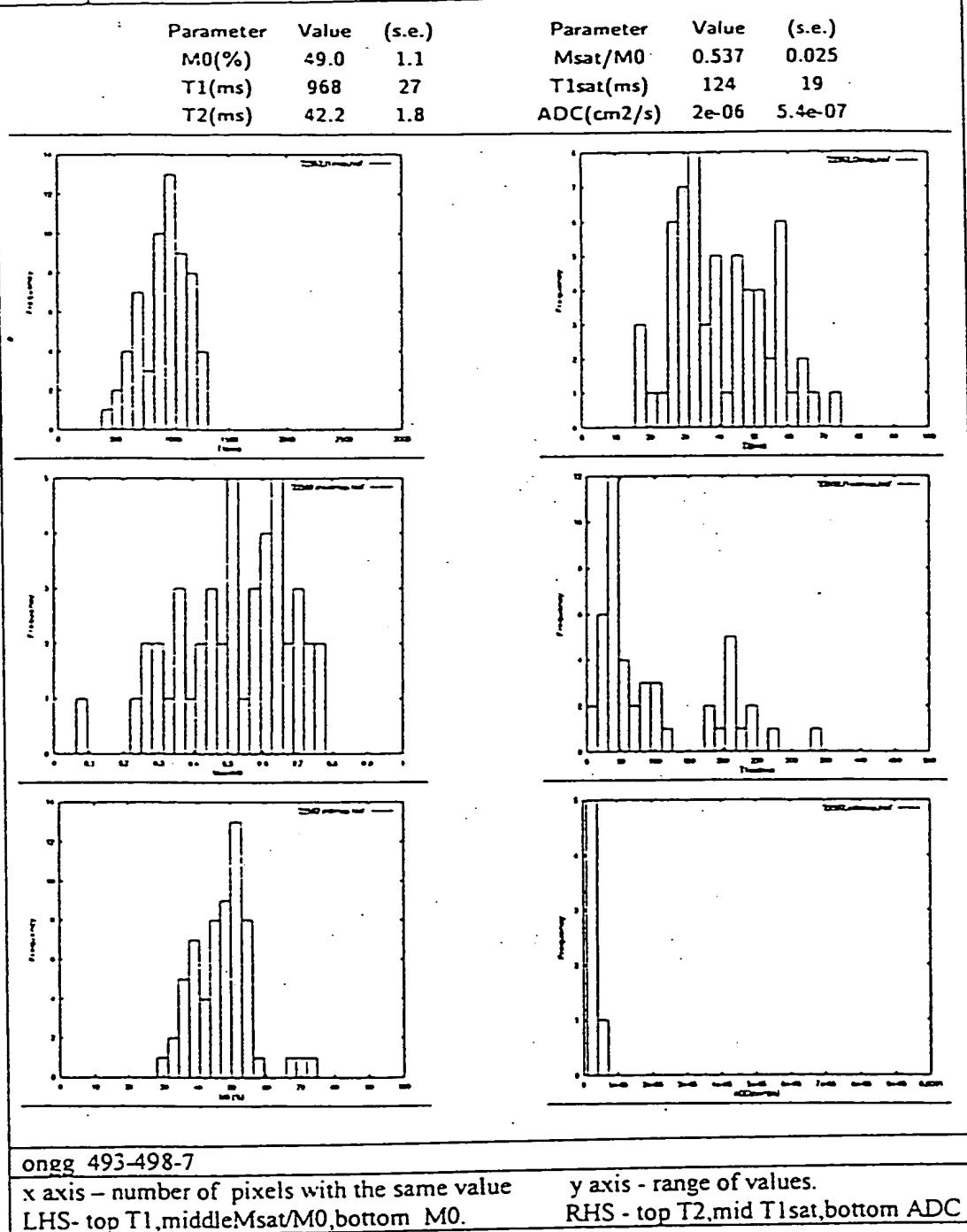


Figure SRL 11. Automated output of histograms for C128 mask 7

C129 (ACD1 + HAED, vital flap) 8 weeks

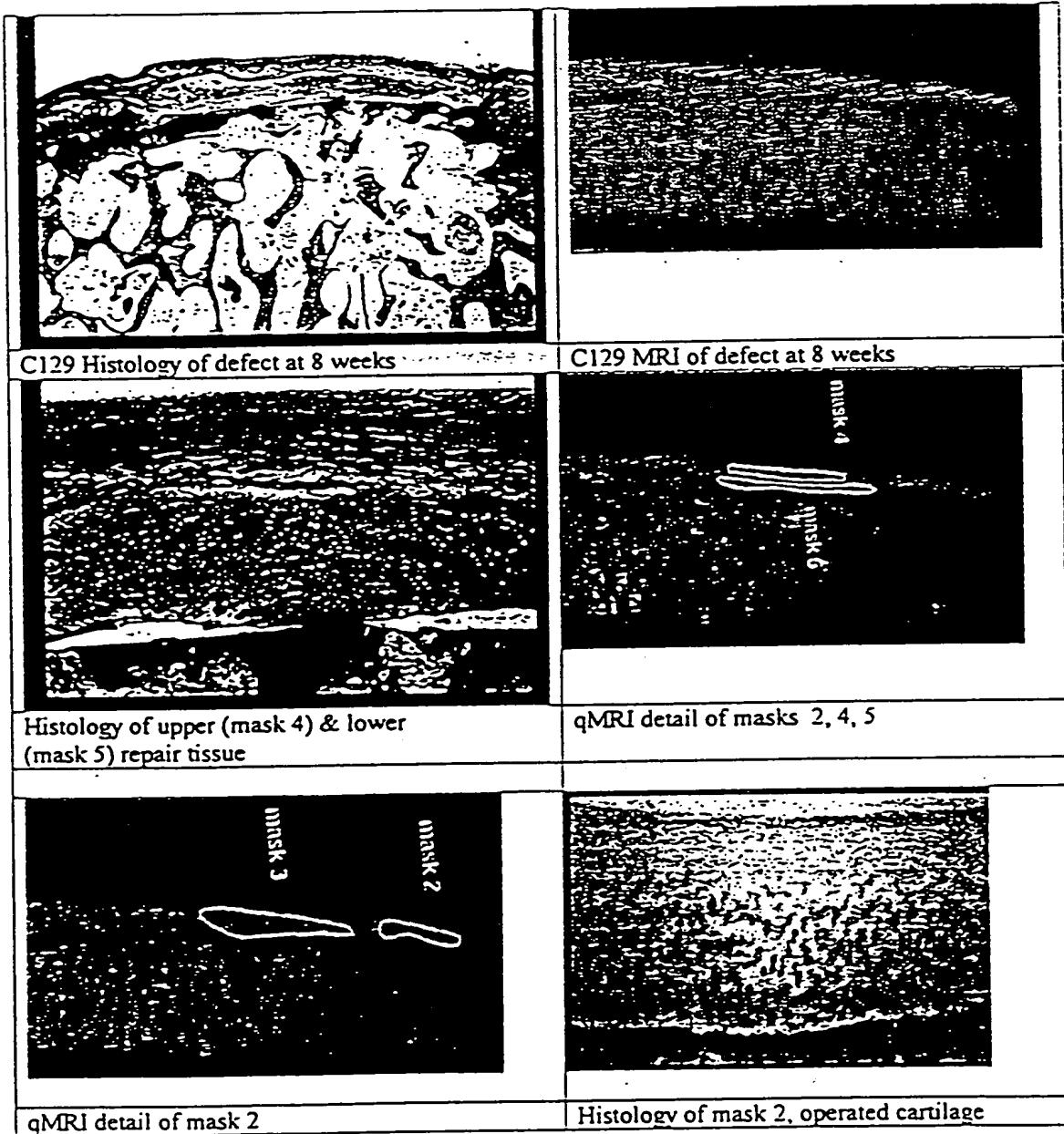


Figure SRL 12.

C129 (ACD1 + HAED, vital flap) 8 weeks

Histology and MRI showing detail of masks 2, 3, 4 and 5.

## C129 qMRI values for mask 2, cartilage adjacent to the defect

Parameter

M0(%)

T1(ms)

T2(ms)

Parameter

Msat/M0

T1sat(ms)

ADC(cm<sup>2</sup>/s)

Value

36.8

1092

14.2

(s.e.)

3.2

173

0.4

Value

0.032

11

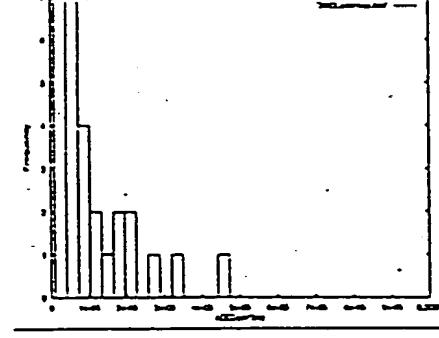
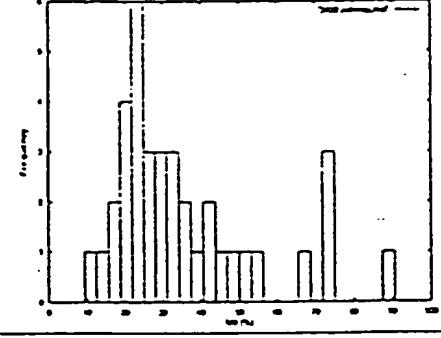
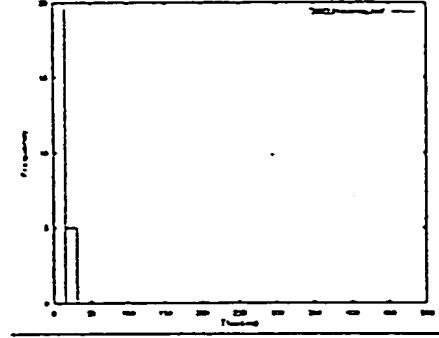
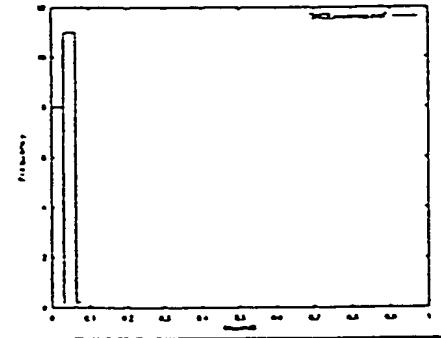
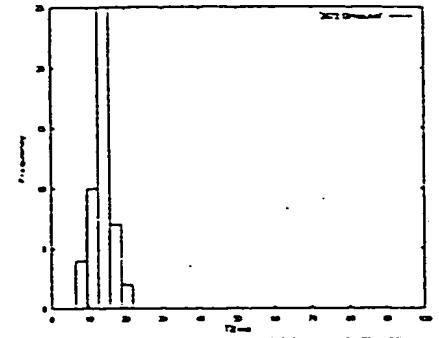
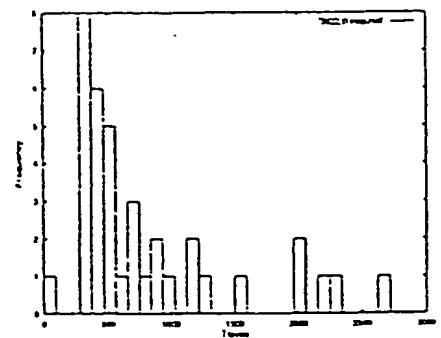
1.1e-05

(s.e.)

0.003

2

2e-06



ongg752-757-2

x axis - number of pixels with the same value  
LHS- top T1, middle Msat/M0, bottom M0.

y axis - range of values.

RHS - top T2, mid T1sat, bottom ADC

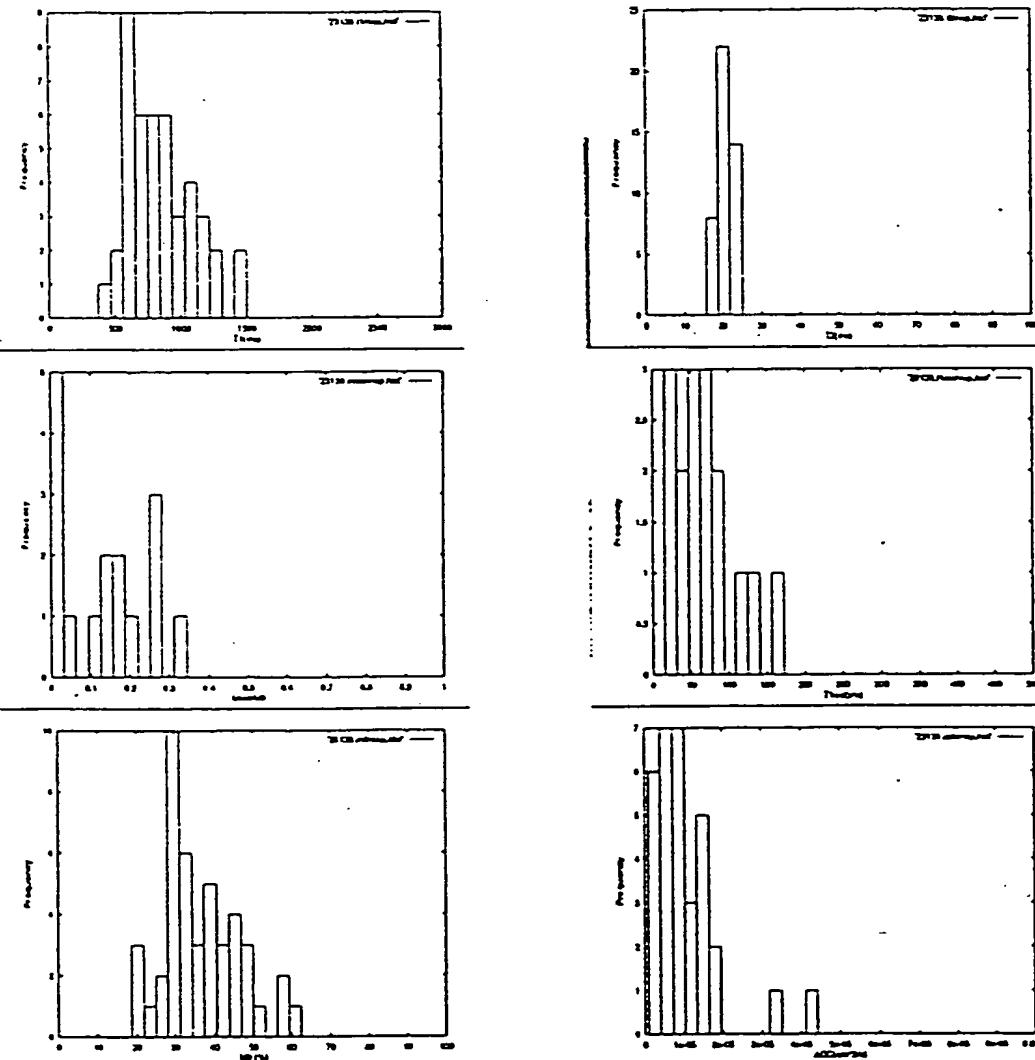
Figure SRL 13. Automated output of histograms for C129 mask 2

105040-02082860

## C129 qMRI values for mask 5, lower repair tissue

Parameter	Value	(s.e.)
M0(%)	38.0	1.6
T1(ms)	876	40
T2(ms)	21.8	0.3

Parameter	Value	(s.e.)
Msat/M0	0.138	0.027
T1sat(ms)	69	11
ADC(cm <sup>2</sup> /s)	1e-05	1.5e-06



ongg752-757-5

x axis - number of pixels with the same value  
 LHS- top T1, middle Msat/M0, bottom M0.

y axis - range of values.  
 RHS - top T2, mid T1sat, bottom ADC

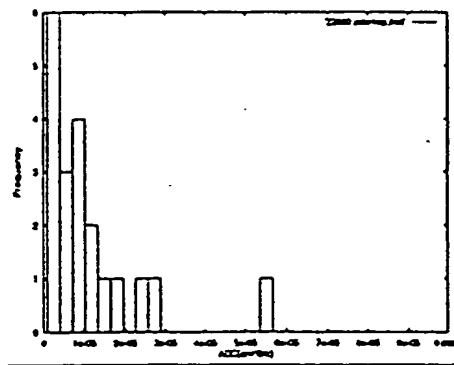
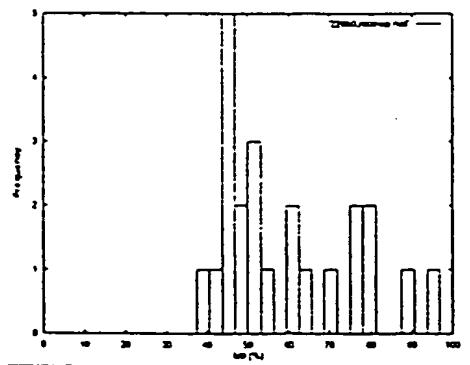
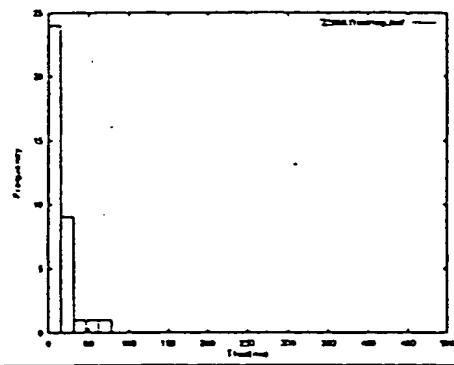
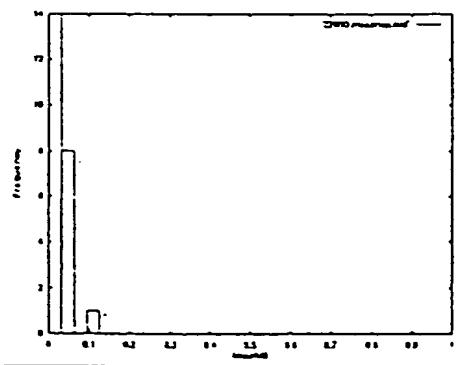
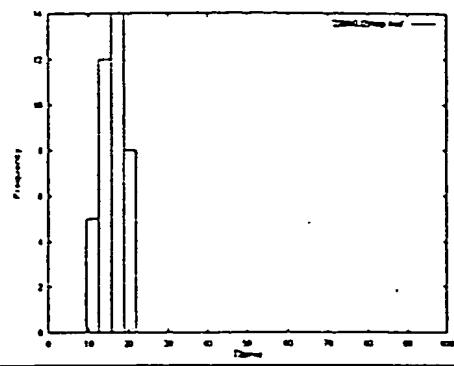
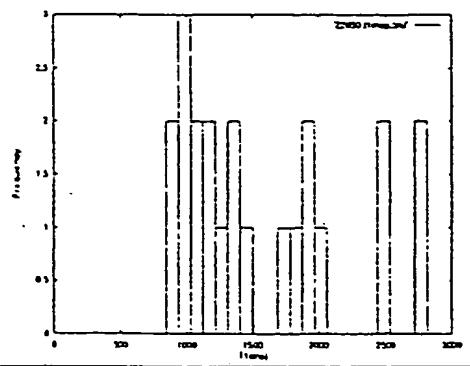
Figure SRL 14. Automated output of histograms for C129 mask 5

00050410-000200000000

## C129 qMRI values for mask 4, upper repair tissue

Parameter	Value	(s.e.)
M0(%)	61.4	3.5
T1(ms)	2025	227
T2(ms)	16.8	0.4

Parameter	Value	(s.e.)
Msat/M0	0.036	0.005
T1sat(ms)	16	3
ADC(cm <sup>2</sup> /s)	1.1e-05	2.7e-06



ongg752-757-4

x axis - number of pixels with the same value  
LHS- top T1, middle Msat/M0, bottom M0.

y axis - range of values.

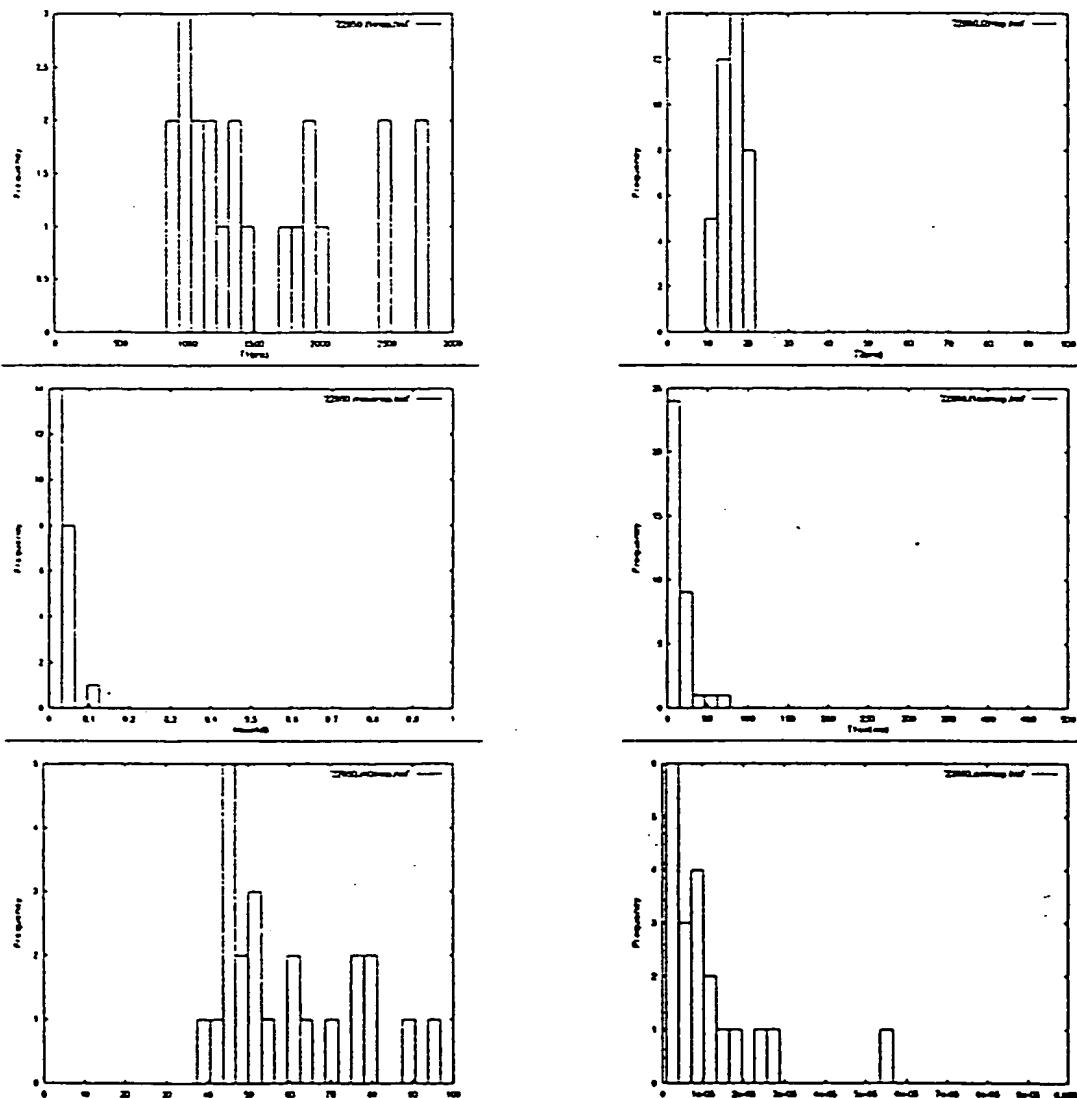
RHS - top T2, mid T1sat, bottom ADC

Figure SRL 15. Automated output of histograms for C129 mask 4

TOS080-020000000000

## C129 qMRI values for mask 3 repair tissue

Parameter	Value	(s.e.)	Parameter	Value	(s.e.)
M0(%)	61.4	3.5	Msat/M0	0.036	0.005
T1(ms)	2025	227	T1sat(ms)	16	3
T2(ms)	16.8	0.4	ADC(cm <sup>2</sup> /s)	1.1e-05	2.7e-06



ongg752-757-3

x axis - number of pixels with the same value  
 LHS- top T1, middle Msat/M0, bottom M0.

y axis - range of values.  
 RHS - top T2, mid T1sat, bottom ADC

Figure SRL 16. Automated output of histograms for C129 mask 3

096280200-00501

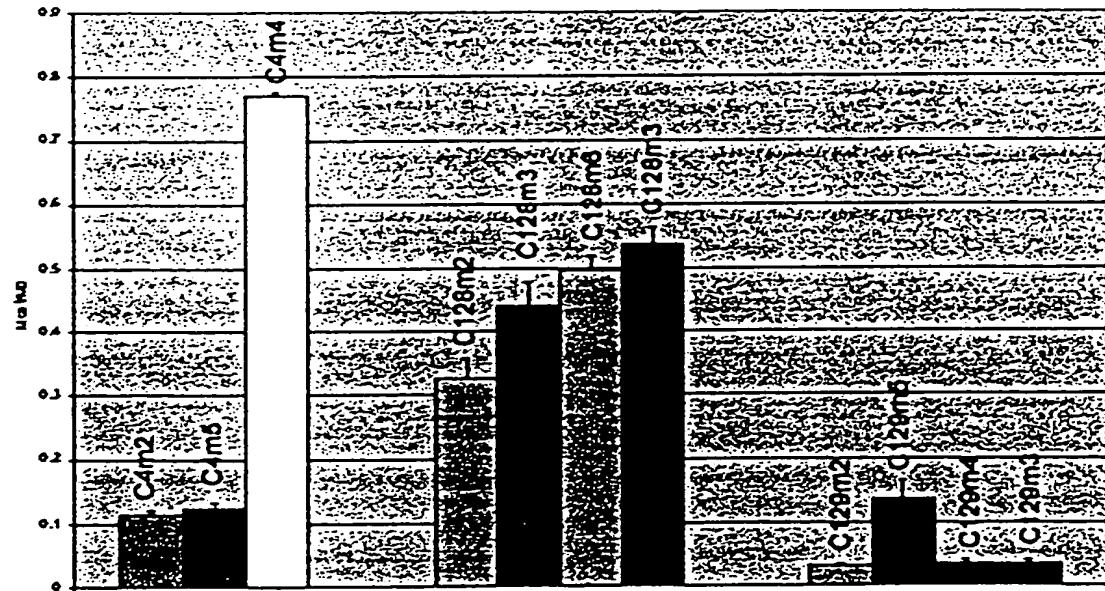
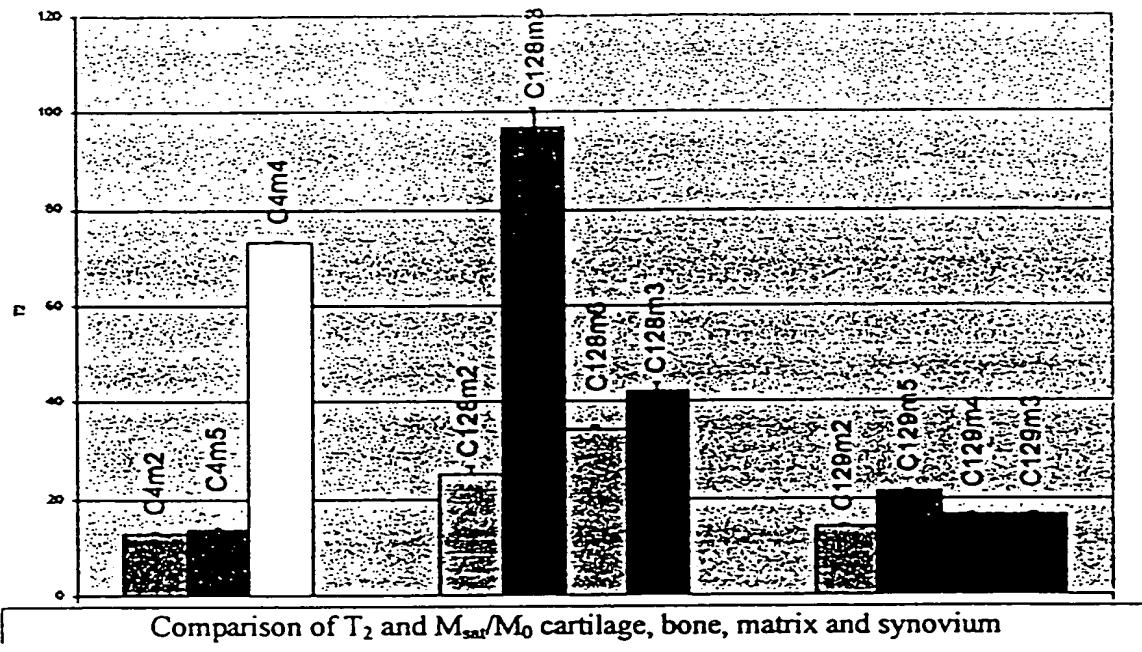
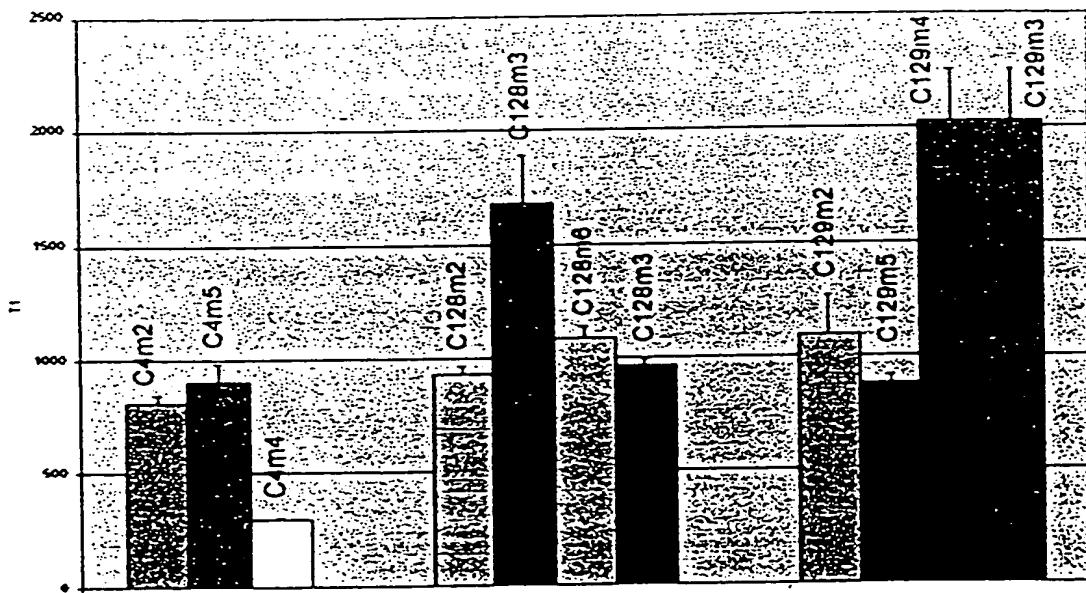
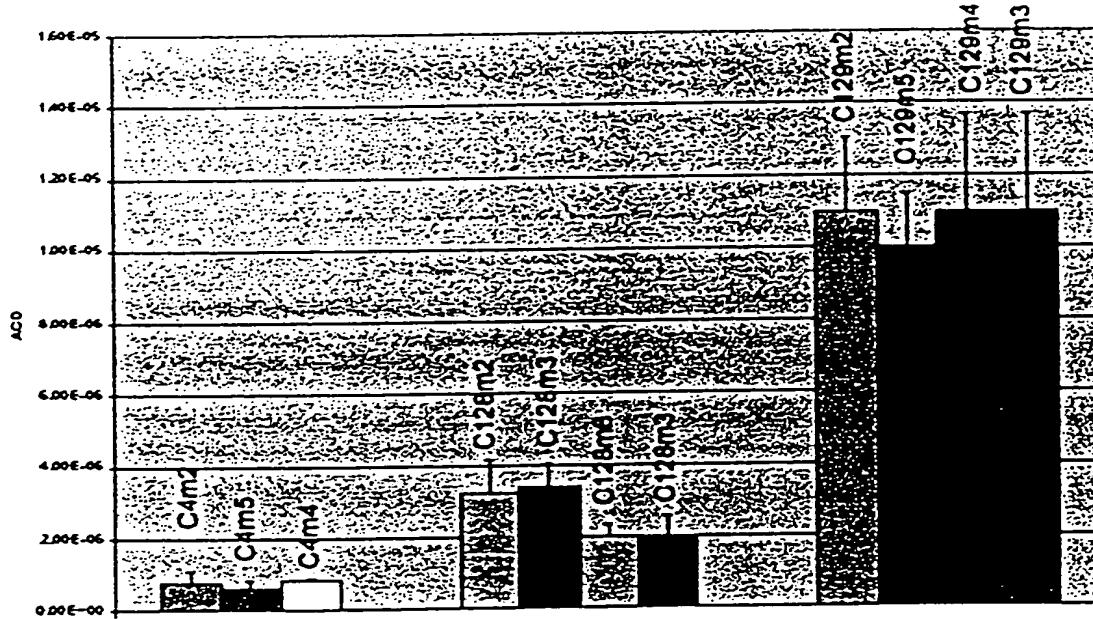


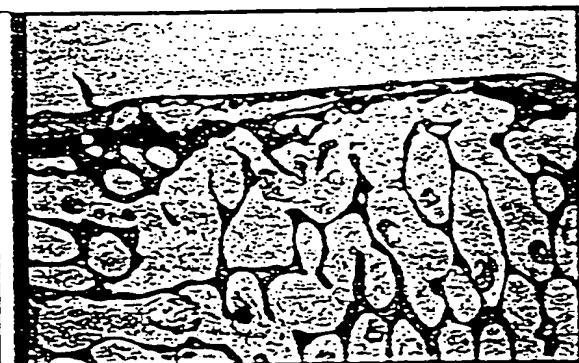
Figure SRL 17a. Graphs comparing mean values of  $T_2$  (above) and  $M_2/M_0$  (below) for ACD1 defects C128 and C129 with the unoperated control, C4

0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9

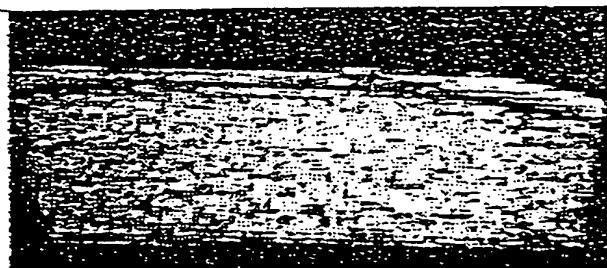
Comparison of T<sub>1</sub> and ADC for cartilage, bone, and synoviumFigure SRL 17b. . Graphs comparing mean values of T<sub>1</sub> (above) and ADC (below) for ACD1 defects C128 and C129 with the unoperated control, C4

T050400-02082360

C146 (ACDI + HAED, vital flap) 8 weeks



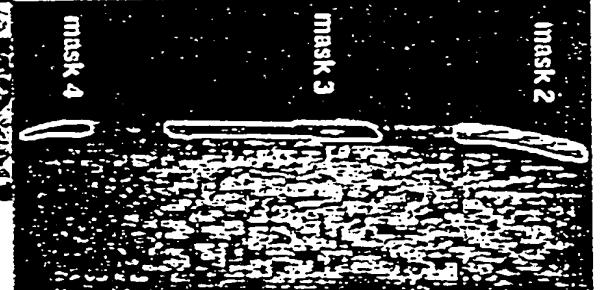
C146 Histology of defect



C146 MRI of defect



Histology detail of mask 3



MRI detail of masks 2, 3, and 4



Histology detail of mask 2

Figure SRL 18

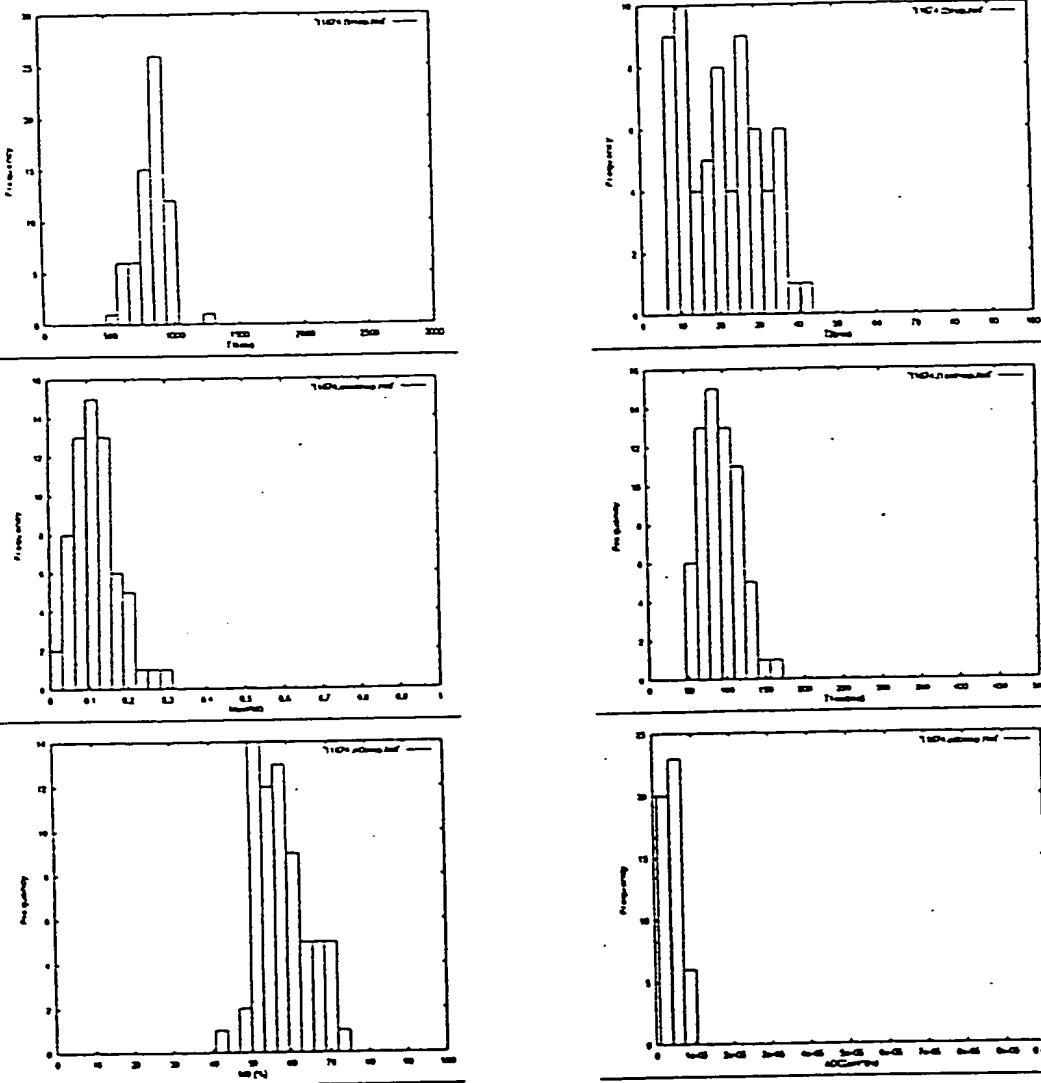
C146 (ACDI + HAED, vital flap) 8 weeks

Histology and MRI showing detail of masks 2, 3 and 4.

09828070 "040501

## C146 qMRI values for mask 2, adjacent cartilage

Parameter	Value	(s.e.)	Parameter	Value	(s.e.)
M0(%)	60.0	0.8	Msat/M0	0.124	0.007
T1(ms)	873	15	T1sat(ms)	97	3
T2(ms)	22.1	1.2	ADC(cm <sup>2</sup> /s)	4.3e-06	3.4e-07



ongg1046-1051-2

x axis - number of pixels with the same value  
LHS- top T1, middle Msat/M0, bottom M0.y axis - range of values.  
RHS - top T2, mid T1sat, bottom ADC

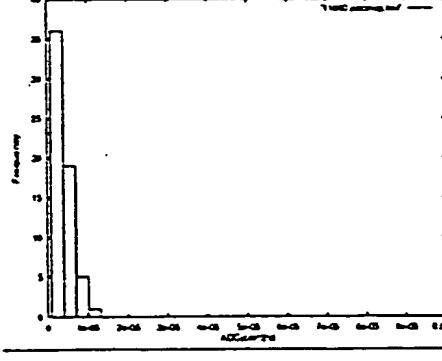
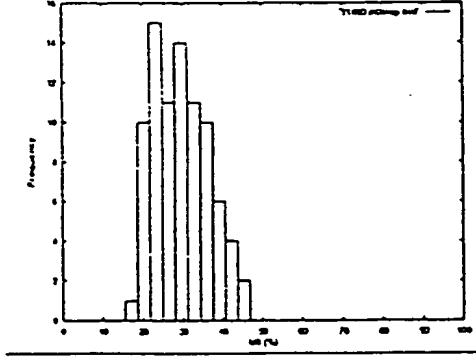
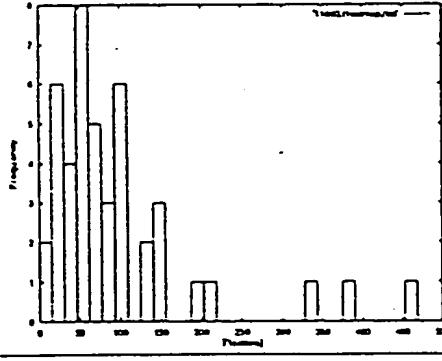
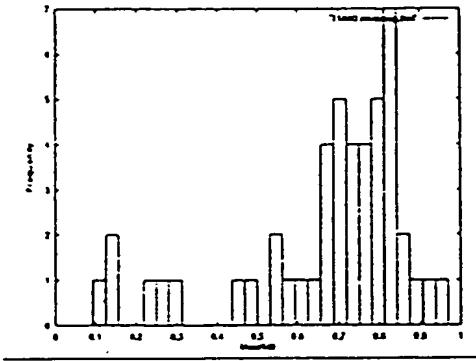
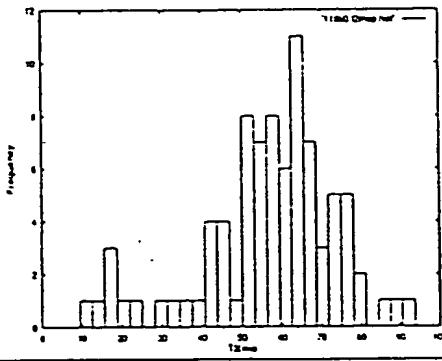
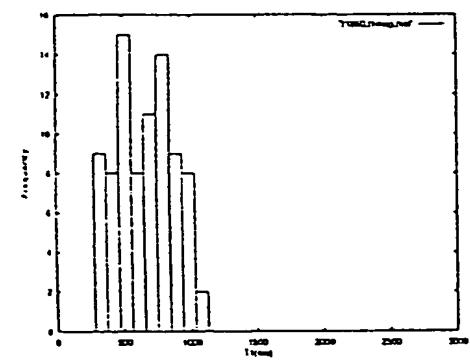
Figure SRL 19. Automated output of histograms for C146 mask 2

098238070 "04/05/09

## C146 qMRI values for mask 3, repair tissue

Parameter	Value	(s.e.)
M0(%)	30.5	0.8
T1(ms)	688	24
T2(ms)	58.9	1.9

Parameter	Value	(s.e.)
Msat/M0	0.692	0.032
T1sat(ms)	119	19
ADC(cm <sup>2</sup> /s)	3.6e-06	3e-07



ongg1046-1051-3

x axis - number of pixels with the same value  
 LHS- top T1, middle Msat/M0, bottom M0.

y axis - range of values.  
 RHS - top T2, mid T1sat, bottom ADC

Figure SRL 20. Automated output of histograms for C146 mask 3

1050400-0200828960

C131 (ACD1 + HAED, partially devitalised flap) 6 weeks

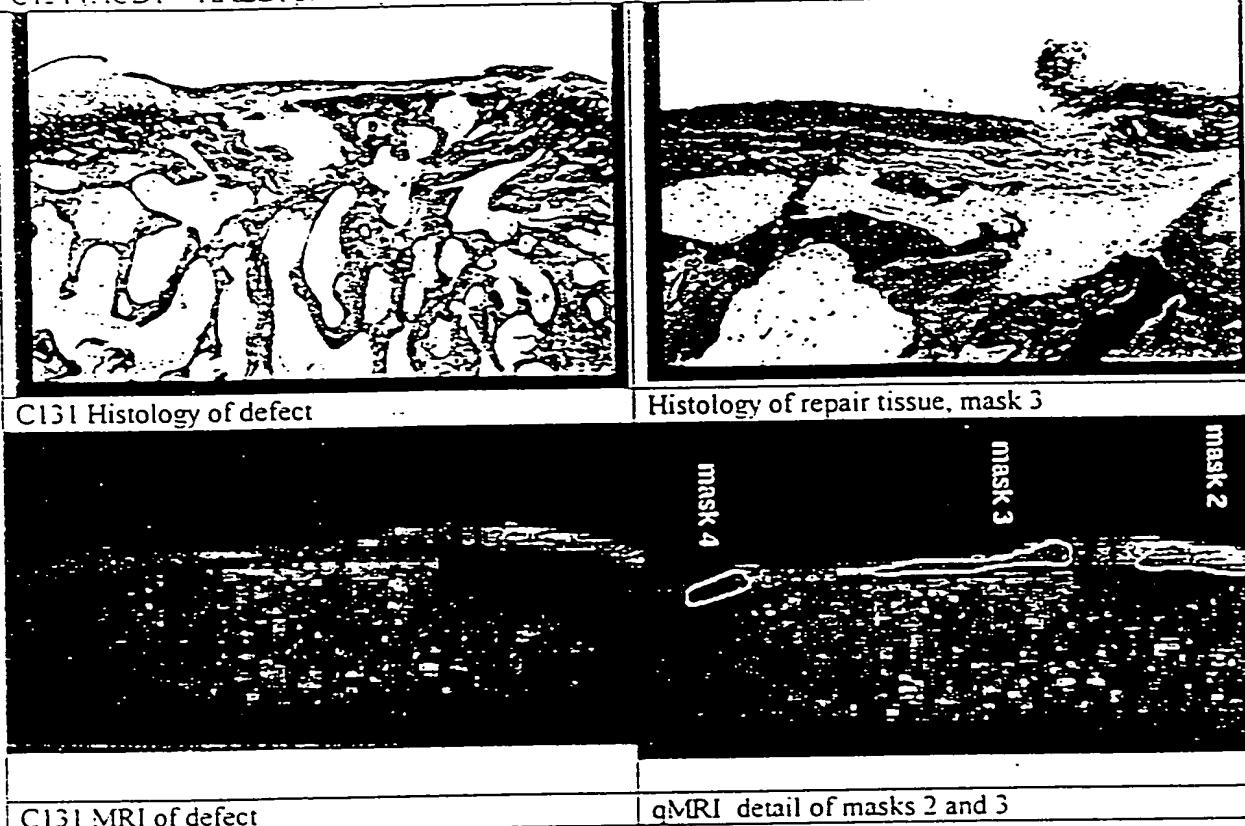
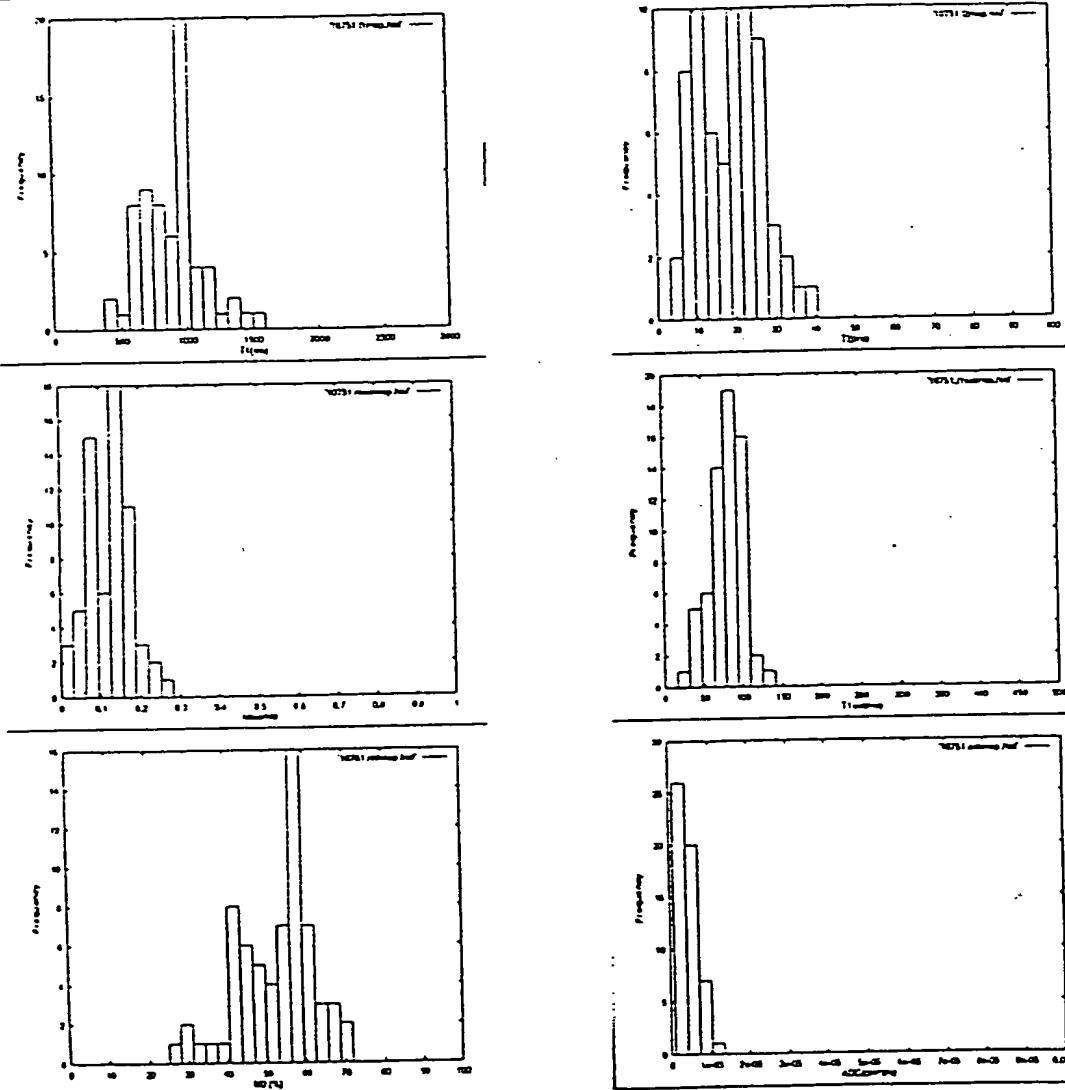


Figure SRL 21  
 C131 (ACDI + HAED, partially devitalised flap) 6 weeks  
 Histology and MRI showing detail of masks 2 and 3.

0 9 8 2 8 0 2 0 0 0 14 0 5 0 4 0

## C131 qMRI values for mask 2, cartilage adjacent to the defect

Parameter	Value	(s.e.)	Parameter	Value	(s.e.)
M0(%)	54.2	1.2	Msat/M0	0.129	0.007
T1(ms)	930	29	T1sat(ms)	84	3
T2(ms)	19.8	1.0	ADC(cm <sup>2</sup> /s)	4.3e-06	3.3e-07



ongg-598-603-2

x axis – number of pixels with the same value  
 LHS- top T1, middle Msat/M0, bottom M0.

y axis - range of values.  
 RHS - top T2, mid T1sat, bottom ADC

Figure SRL 22. Automated output of histograms for C131 mask 2

09828070 "040500

## C131 qMRI values for mask 3, repair tissue

Parameter Value (s.e.)

M0(%) 41.7 1.1

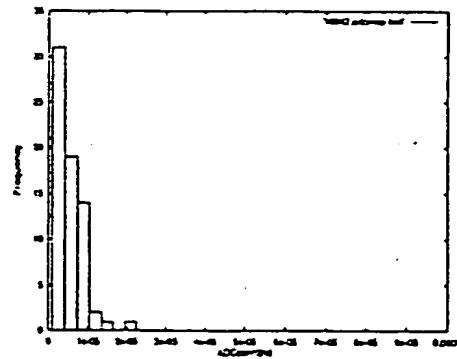
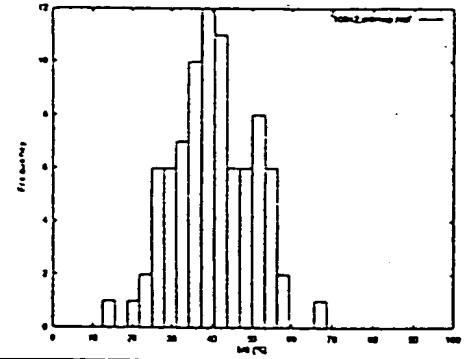
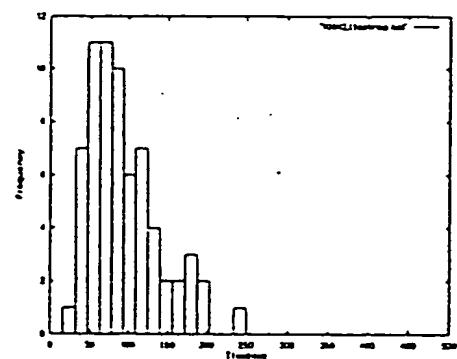
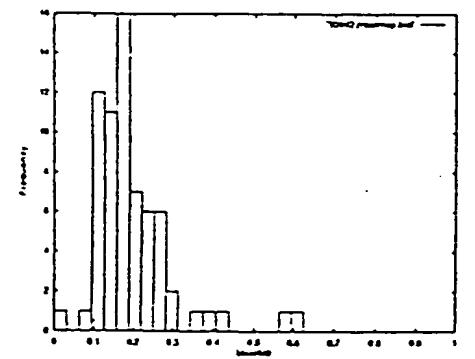
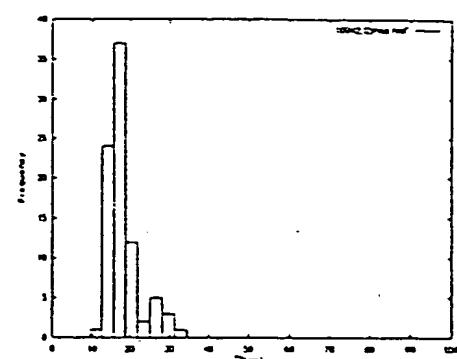
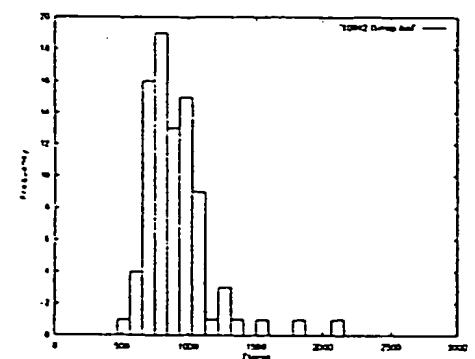
T1(ms) 936 28

T2(ms) 18.6 0.5

Parameter Value (s.e.)

Msat/M0 0.202 0.013

T1sat(ms) 97 6

ADC(cm<sup>2</sup>/s) 5.3e-06 4.3e-07

ongg 598-603-3

x axis - number of pixels with the same value

LHS- top T1, middle Msat/M0, bottom M0.

y axis - range of values.

RHS - top T2, mid T1sat, bottom ADC

Figure SRL 23. Automated output of histograms for C131 mask 3

T050840 "02082860

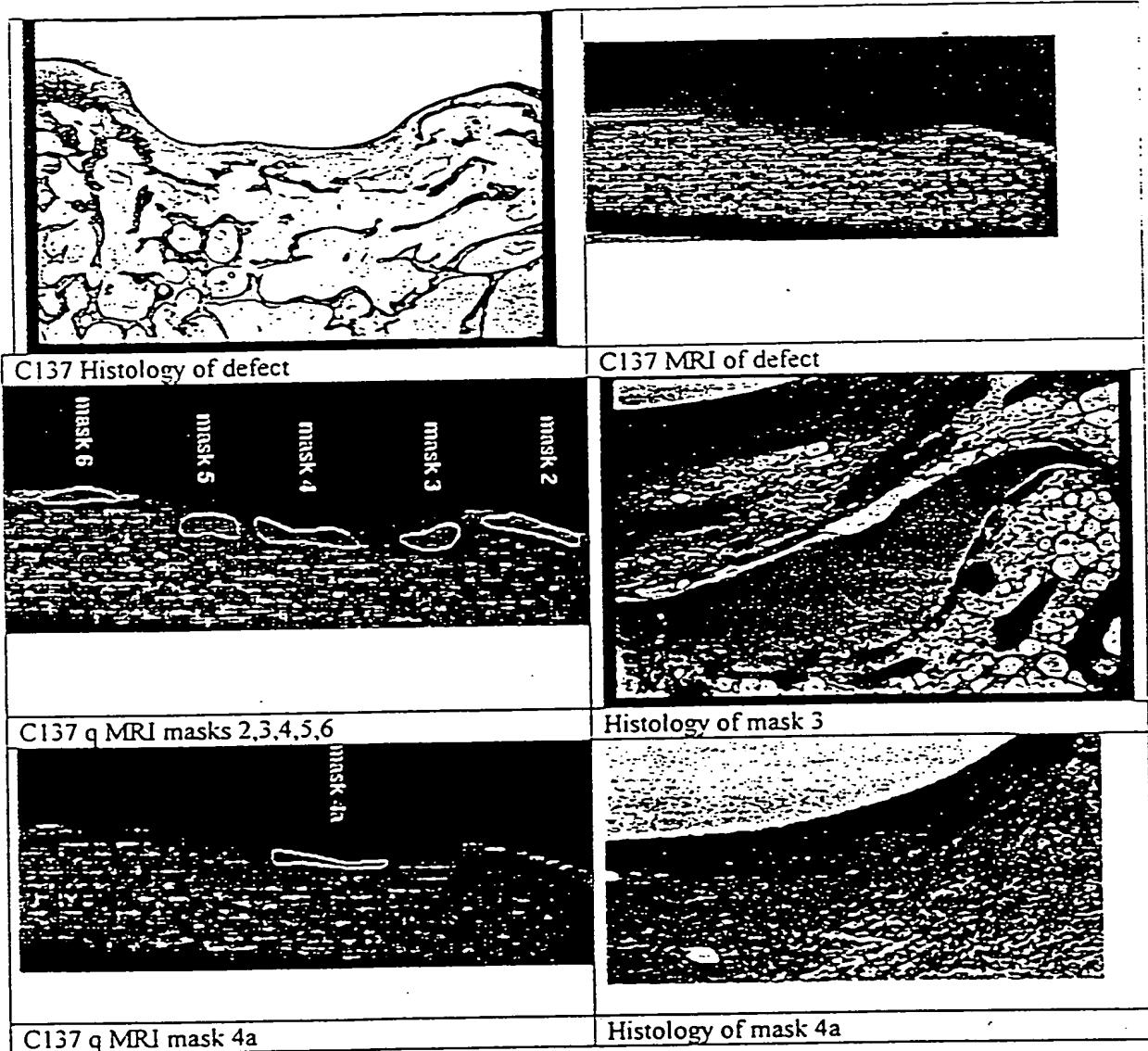


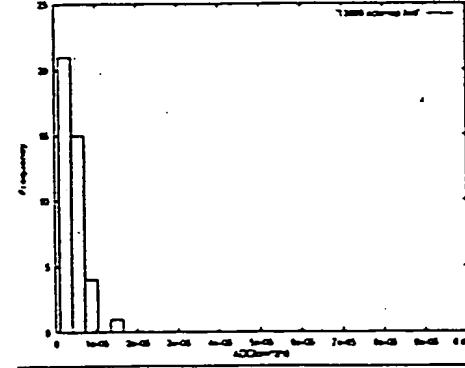
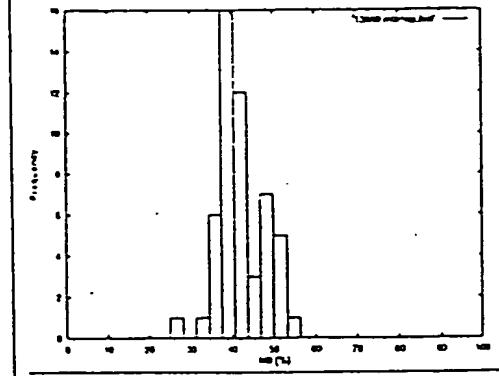
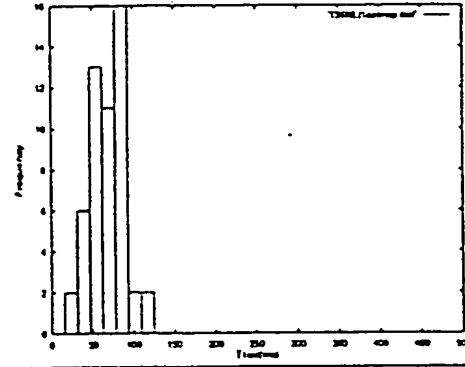
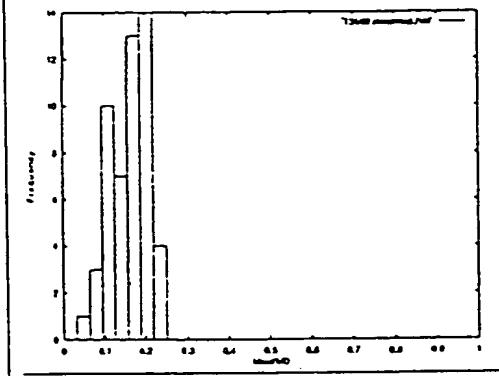
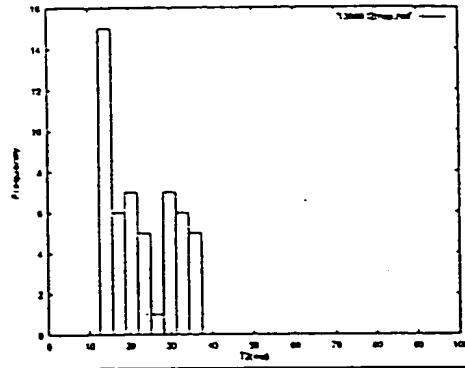
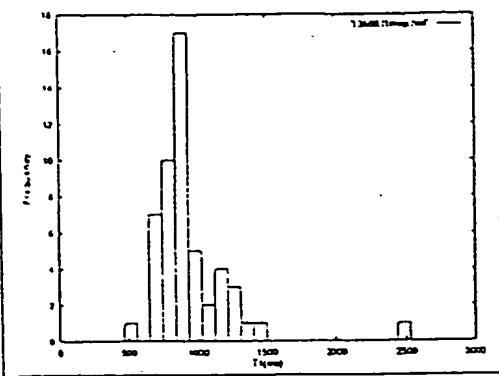
Figure SRL 24.

C137 (virtual ACD1 + PEG Triacrylate, partially devitalised flap) 10 weeks

Histology and MRI showing detail of masks 2, 3, 4, 4a, 5 and 6.

C137 mask 2, cartilage adjacent to the defect

Parameter	Value	(s.e.)	Parameter	Value	(s.e.)
M0(%)	43.6	0.8	Msat/M0	0.167	0.007
T1(ms)	982	40	T1sat(ms)	72	3
T2(ms)	23.6	1.1	ADC(cm <sup>2</sup> /s)	4.3e-06	4.2e-07



ongg 941-946-2

x axis - number of pixels with the same value  
 LHS- top T1, middle  $M_{sat}/M_0$ , bottom  $M_0$ .

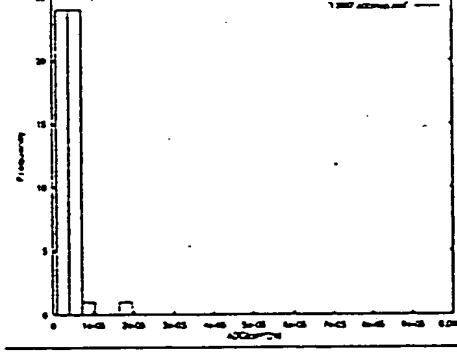
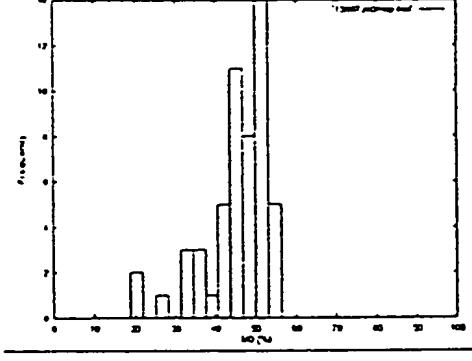
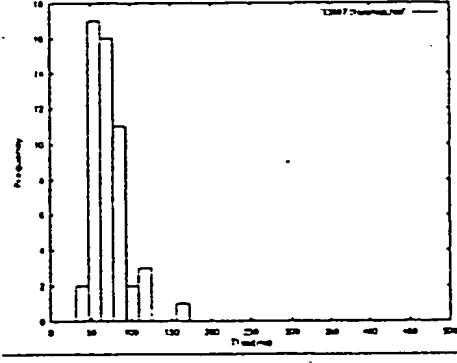
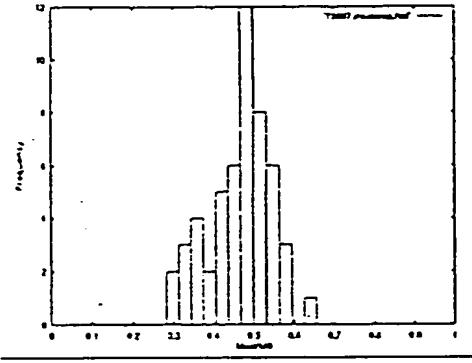
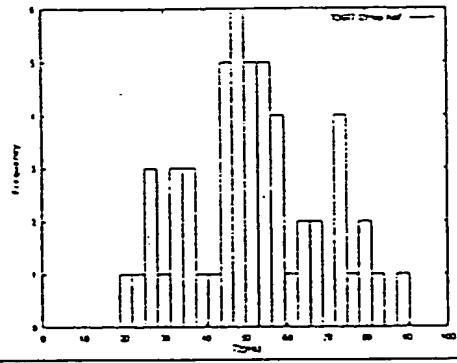
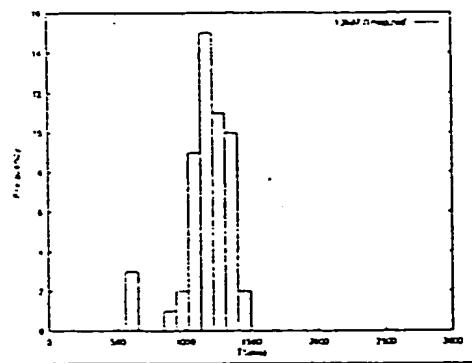
y axis - range of values.

RHS - top T2, mid T1 sat, bottom ADC

Figure SRL 25. Automated output of histograms for C137 mask 2

### C137 mask 3, retained matrix

Parameter	Value	(s.e.)	Parameter	Value	(s.e.)
M0(%)	46.8	1.1	Msat/M0	0.430	0.011
T1(ms)	1219	26	T1sat(ms)	75	3
T2(ms)	53.4	2.3	ADC(cm <sup>2</sup> /s)	4.5e-06	3.7e-07



ongg 941-946-3

x axis – number of pixels with the same value  
LHS- top T1, middle  $M_{sat}/M_0$ , bottom  $M_0$ .

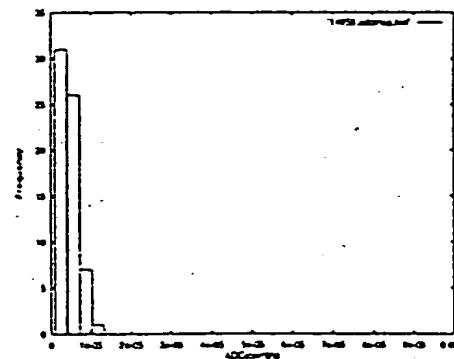
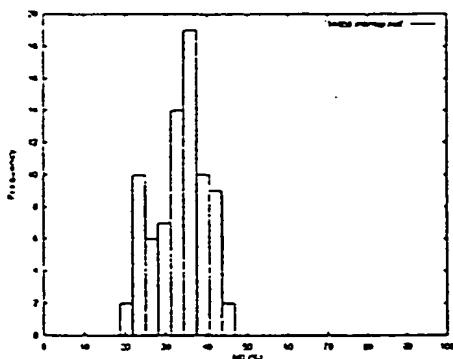
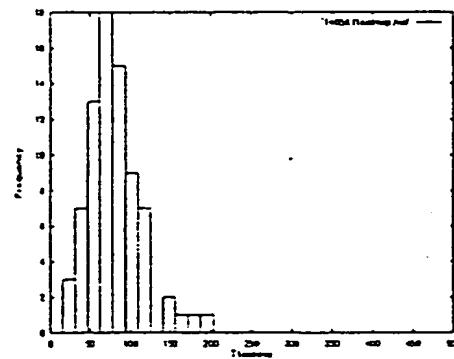
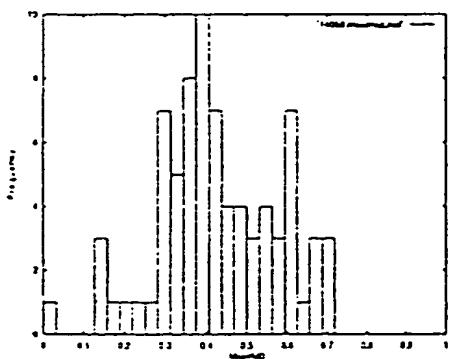
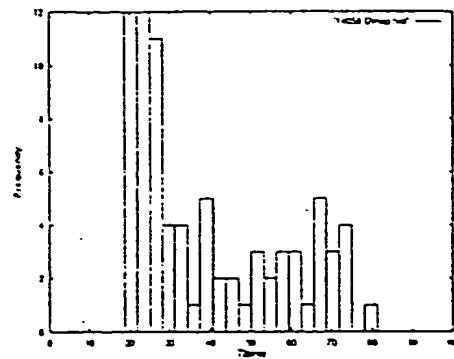
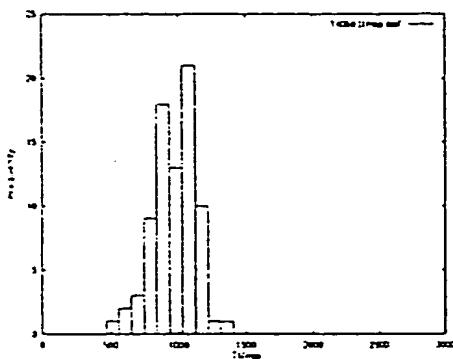
y axis - range of values.

RHS - top T2, mid T1 sat, bottom ADC

Figure SRL 26. Automated output of histograms for C137 mask 3

### C137 mask 4, lower repair tissue

Parameter	Value	(s.e.)	Parameter	Value	(s.e.)
M0(%)	34.8	0.7	Msat/M0	0.443	0.017
T1(ms)	1009	18	T1sat(ms)	83	4
T2(ms)	40.9	2.2	ADC(cm <sup>2</sup> /s)	4.4e-06	3e-07



ongg 941-946-4

x axis - number of pixels with the same value  
 LHS- top T1,middleMsat/M0, bottom M0.

y axis - range of values.

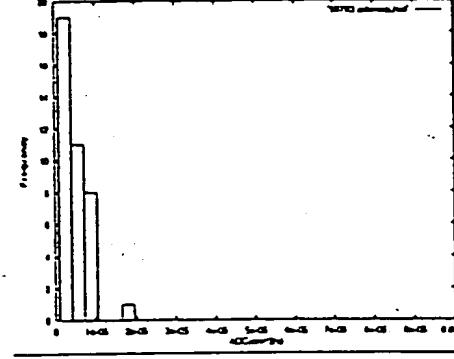
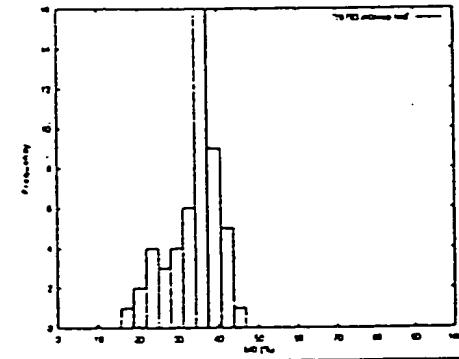
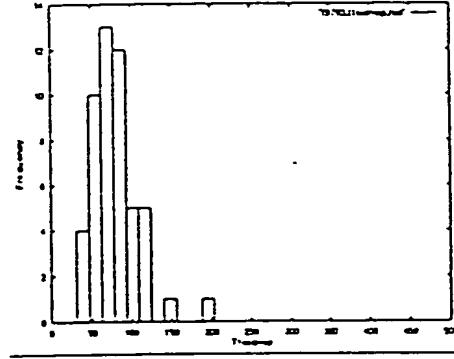
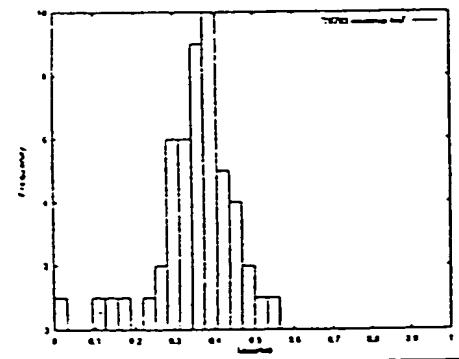
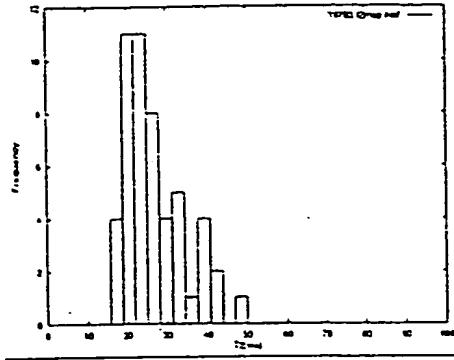
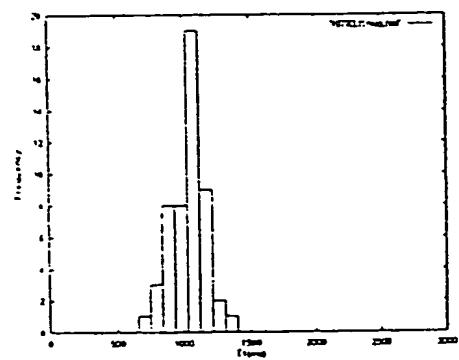
RHS - top T2,mid T1 sat, bottom ADC

Figure SRL 27. Automated output of histograms for C137 mask 4

## C137 mask 4a, upper repair tissue

Parameter	Value	(s.e.)
M0(%)	35.4	0.9
T1(ms)	1080	19
T2(ms)	27.7	1.1

Parameter	Value	(s.e.)
Msat/M0	0.365	0.014
T1sat(ms)	83	4
ADC(cm <sup>2</sup> /s)	5e-06	5.2e-07



ongg 941-946-4a

x axis - number of pixels with the same value  
LHS- top T1, middle Msat/M0, bottom M0.

y axis - range of values.

RHS - top T2, mid T1sat, bottom ADC

Figure SRL 28. Automated output of histograms for C137 mask 4a

00000000000000000000000000000000

## C137 mask 5, retained matrix

Parameter

Value

(s.e.)

M0(%)

42.7

1.3

T1(ms)

1173

42

T2(ms)

56.4

1.7

Parameter

Value

(s.e.)

Msat/M0

0.576

0.014

T1sat(ms)

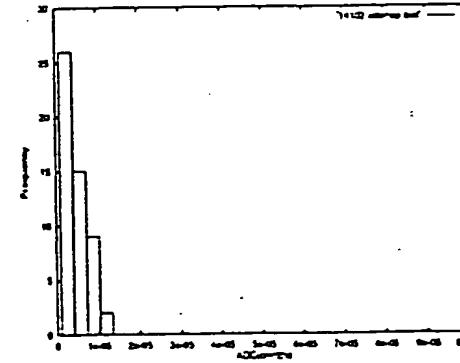
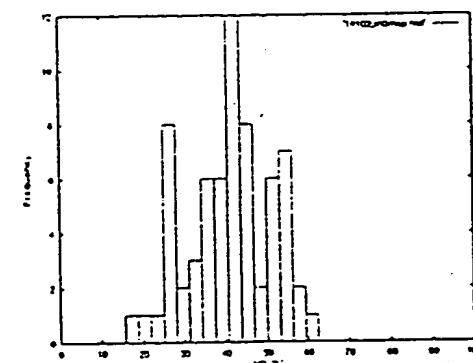
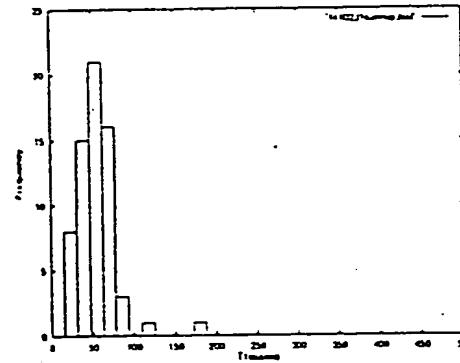
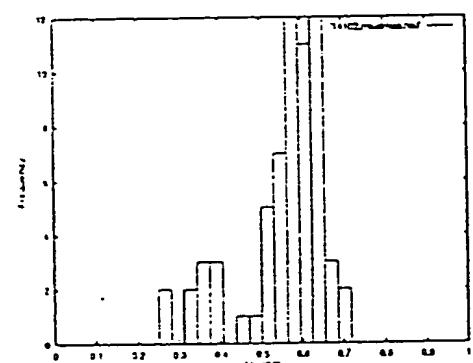
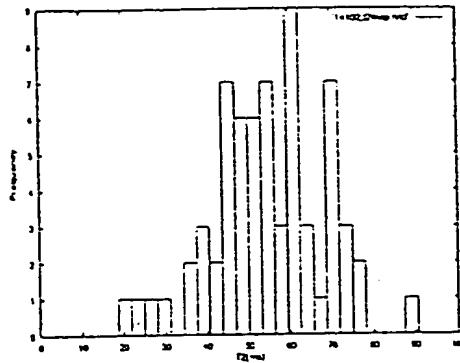
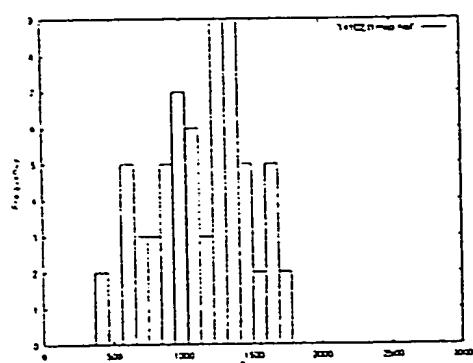
57

3

ADC(cm<sup>2</sup>/s)

4.5e-06

3.7e-07



0000941-946-5

x axis - number of pixels with the same value

y axis - range of values.

LHS- top T1, middle Msat/M0, bottom M0.

RHS - top T2, mid T1sat, bottom ADC

Figure SRL 29. Automated output of histograms for C137 mask 5

0000941-946-5

qMRI analysis of repair tissue in ACD3 defects

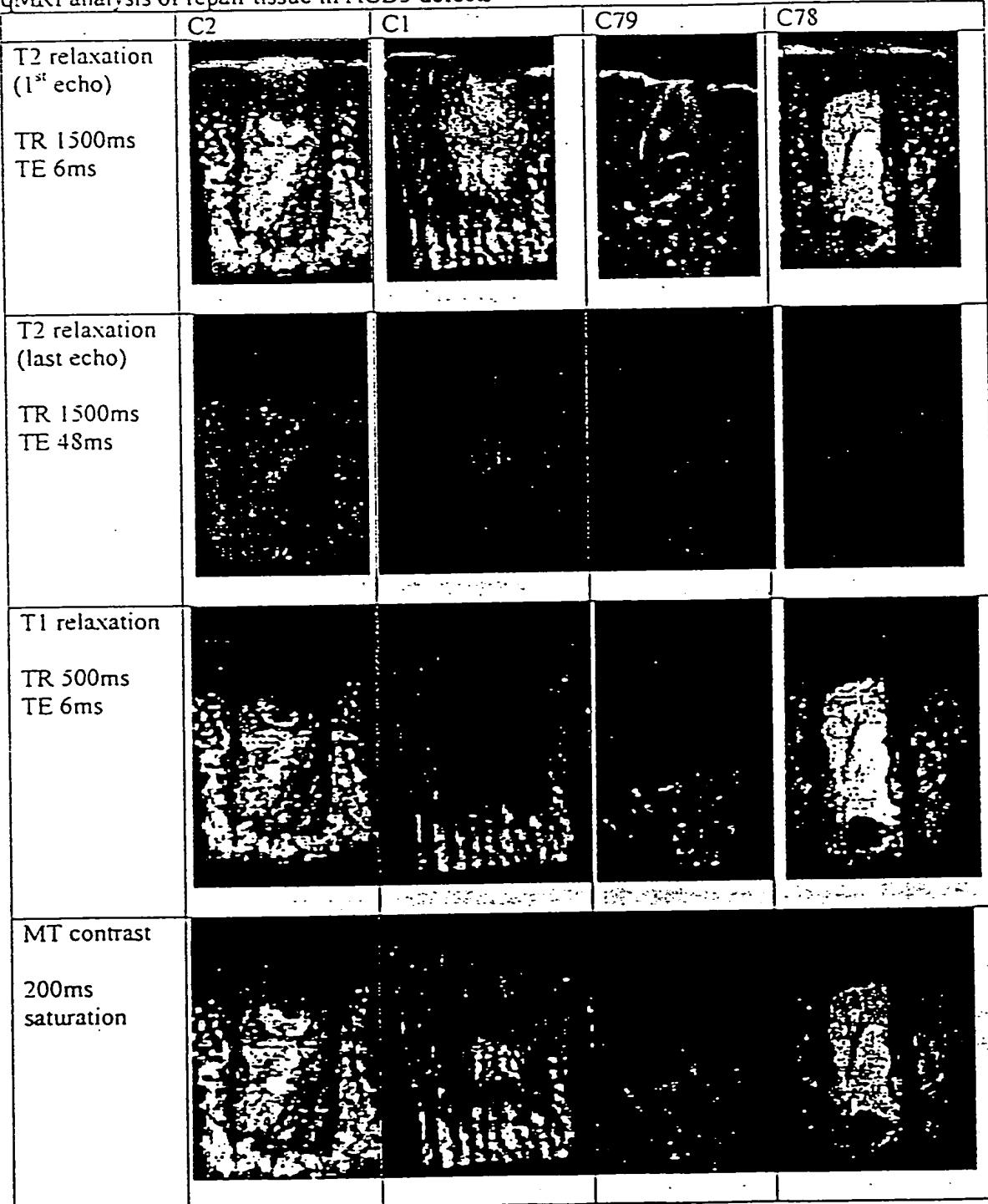


Figure SRL 30.

Selected MR scans from qMRI analysis of spontaneously healed ACD3 defects  
C1, C2 at 8 weeks and C78, C79 at 6 months.

P050400-02082860

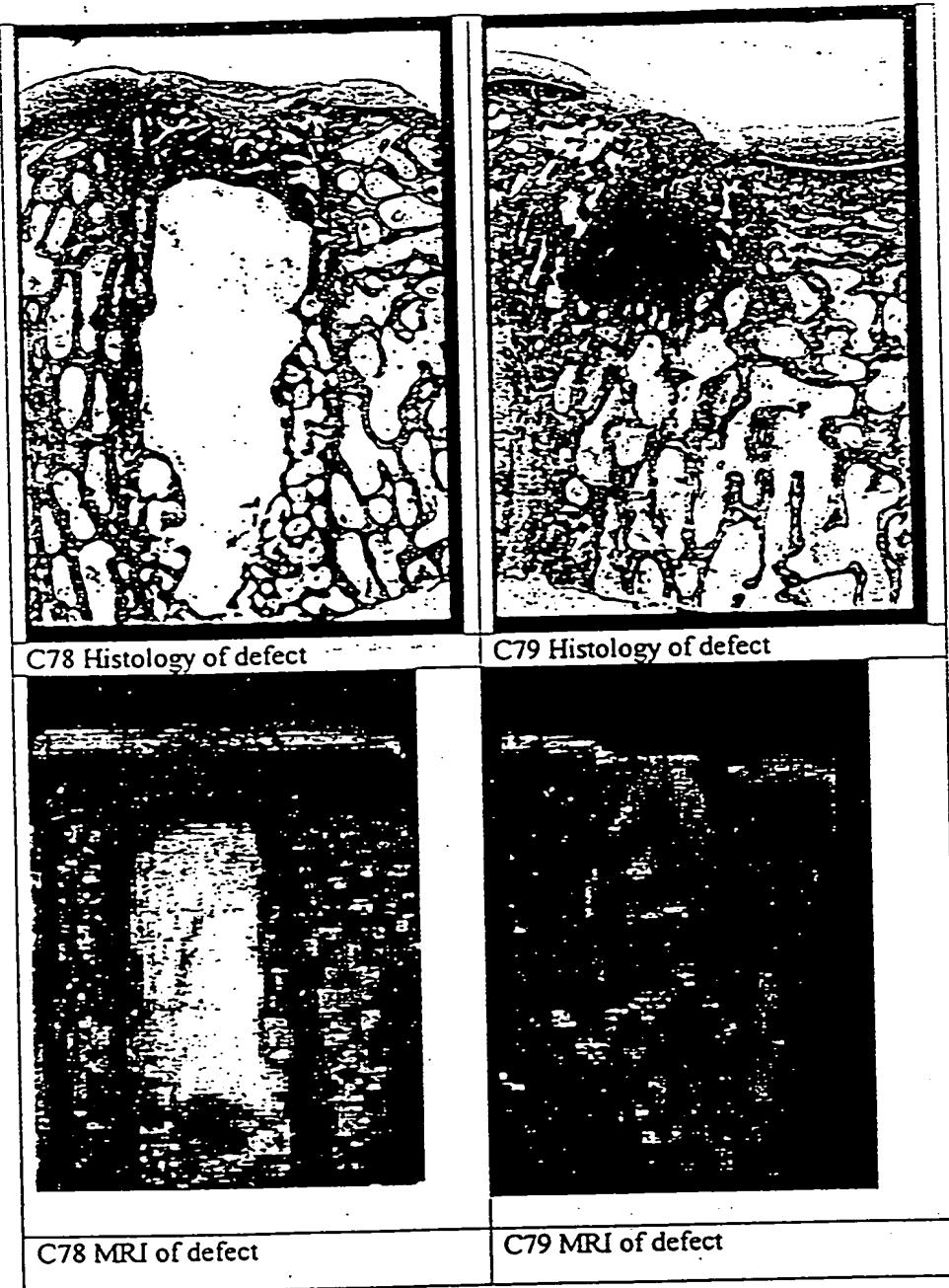


Figure SRL 31  
Histology and MRI of spontaneously healed ACD3 defects C78 and  
C79 at 6 months

0004070008828260

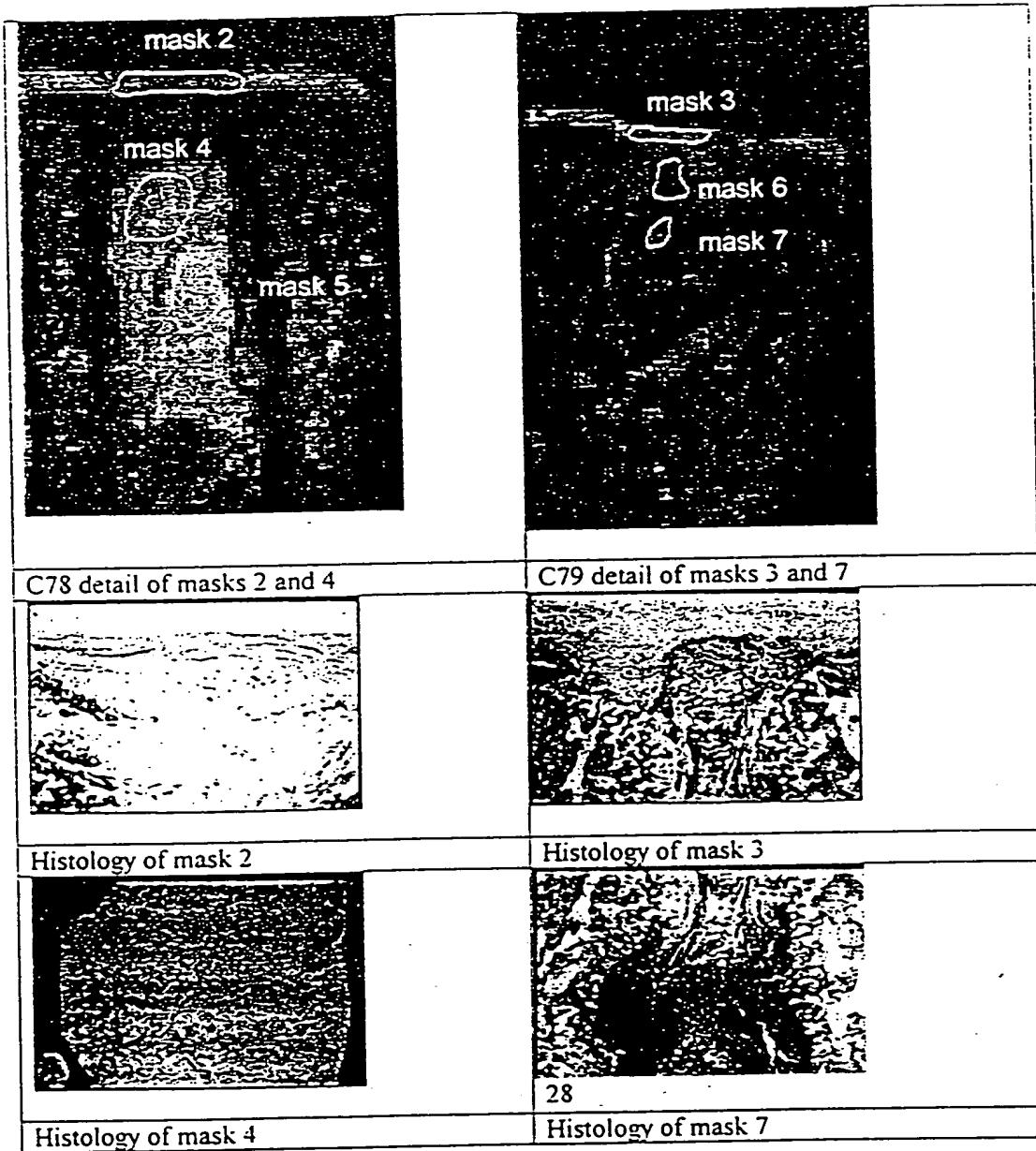
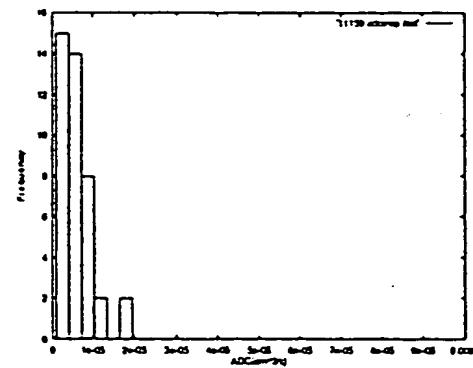
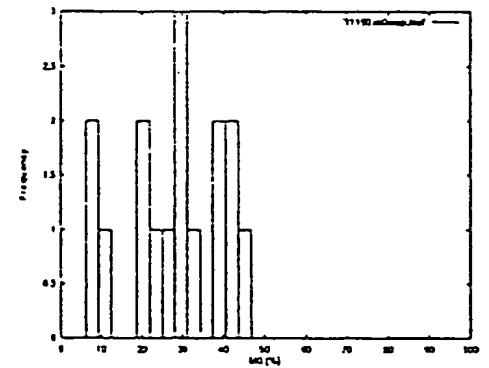
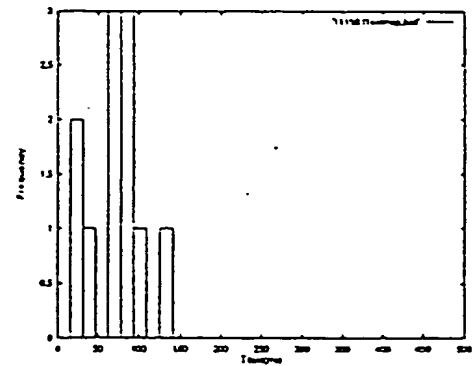
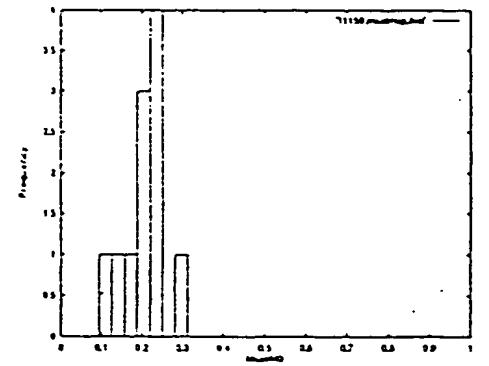
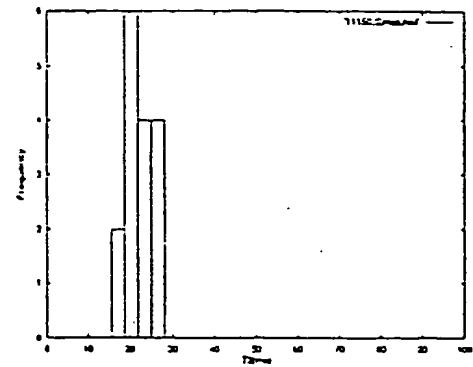
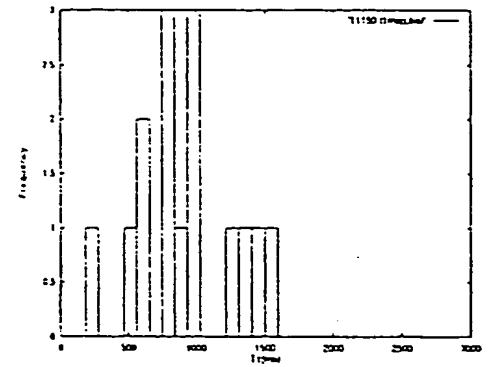


Figure SRL 32  
Histology and MRI showing detail of masks 2, 4, 3 and 7 for C78 and C79

C78 qMRI values for mask 2, repair tissue in cartilage compartment

Parameter	Value	(s.e.)	Parameter	Value	(s.e.)
M0(%)	28.6	3.2	Msat/M0	0.215	0.016
T1(ms)	1095	159	T1sat(ms)	75	10
T2(ms)	23.0	0.8	ADC(cm <sup>2</sup> /s)	1.4e-05	8.1e-06



ongg 0004-0009-2

x axis – number of pixels with the same value  
LHS- top T1, middle  $M_{sat}/M_0$ , bottom  $M_0$ .

y axis - range of values.

RHS top T2,mid T1sat, bottom ADC

Figure SRL 33 Automated output of histograms for C78 mask 2

C78 qMRI values for mask 4, repair tissue in bone compartment

Parameter Value (s.e.)

M0(%) 24.0 2.6

T1(ms) 1034 68

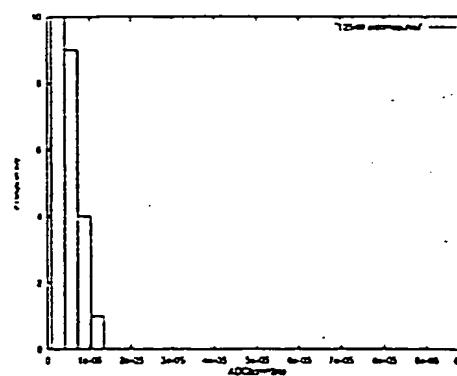
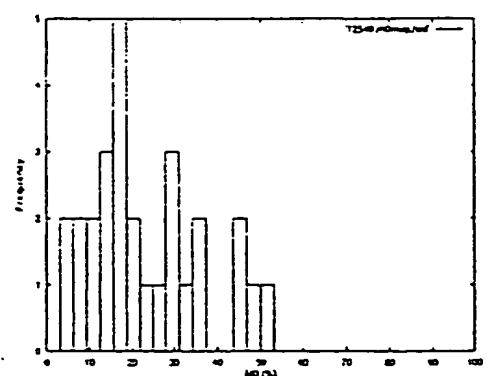
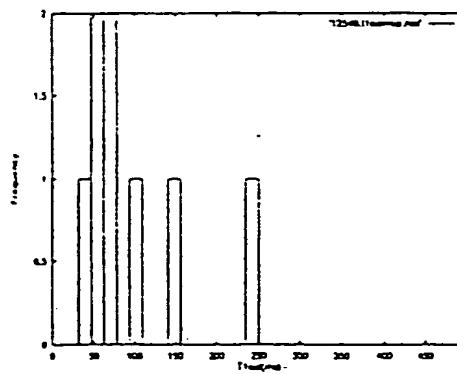
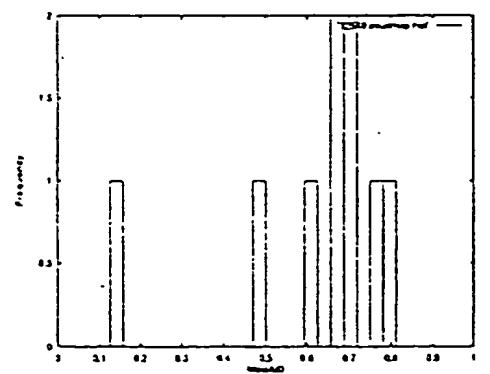
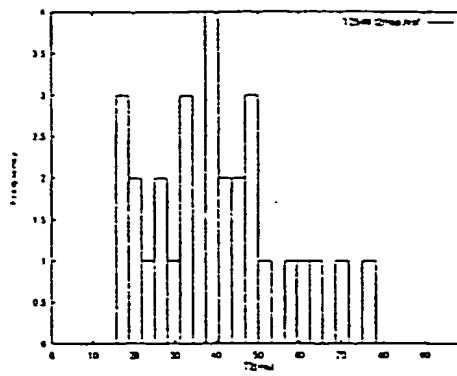
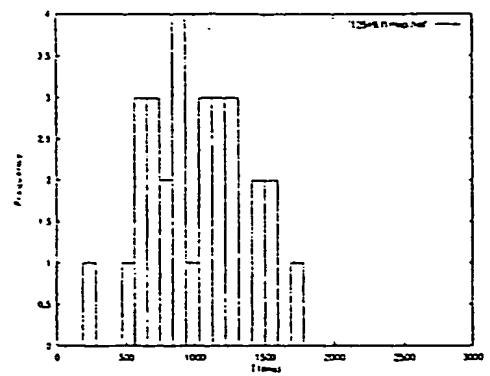
T2(ms) 41.0 3.1

Parameter Value (s.e.)

Msat/M0 0.641 0.069

T1sat(ms) 101 24

ADC(cm<sup>2</sup>/s) 5.1e-06 5.8e-07



ongg 0004-0009-4

x axis - number of pixels with the same value  
 LHS- top T1,middleMsat/M0,bottom M0.

y axis - range of values.

RHS top T2,mid T1sat,bottom ADC

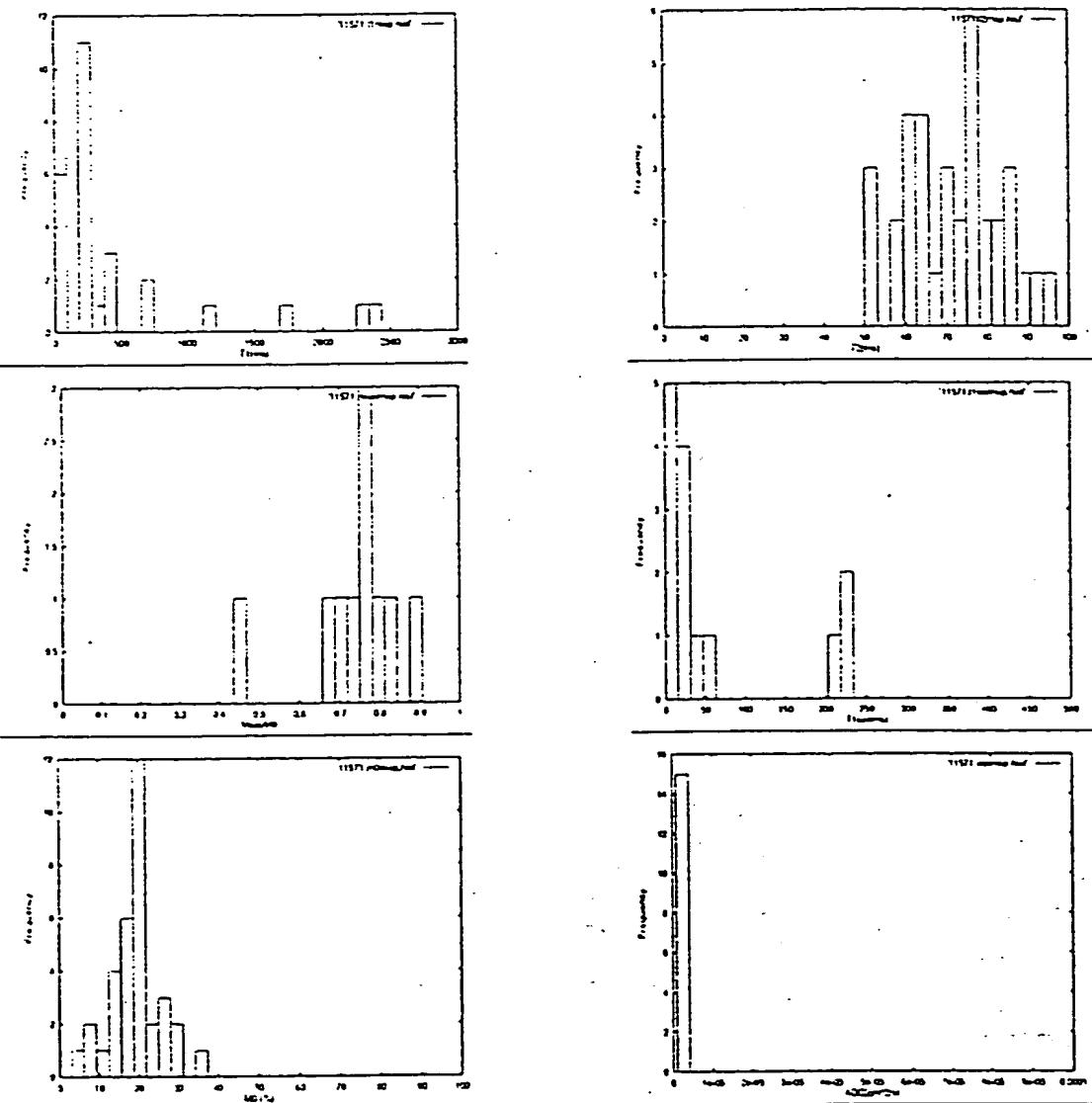
Figure SRL 34 Automated output of histograms for C78 mask 4

105040402082860

## C79 qMRI values for mask 3, repair tissue in cartilage compartment

Parameter	Value	(s.e.)
M0(%)	20.1	1.2
T1(ms)	427	106
T2(ms)	75.8	2.6

Parameter	Value	(s.e.)
Msat/M0	0.763	0.038
T1sat(ms)	65	24
ADC( $\text{cm}^2/\text{s}$ )	1.5e-06	1.9e-07



long 0004-0009-3

x axis - number of pixels with the same value  
LHS - top T1, middle Msat/M0, bottom M0.

y axis - range of values.

RHS top T2, mid T1sat, bottom ADC

Figure SRL 35 Automated output of histograms for C79 mask 3

FD5040-020000000000

## C79 qMRI values for mask 7, repair tissue in bone compartment

## Parameter

M0(%)

T1(ms)

T2(ms)

(s.e.)

13.5

125

2.8

## Parameter

Msat/M0

T1sat(ms)

ADC(cm<sup>2</sup>/s)

Value

0.674

62

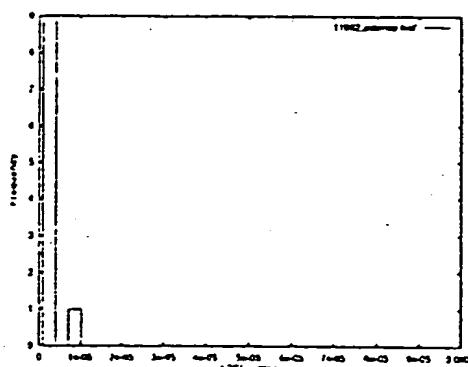
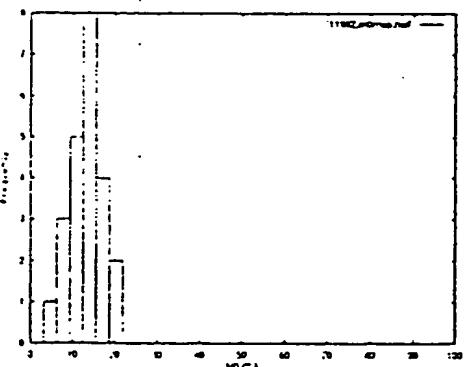
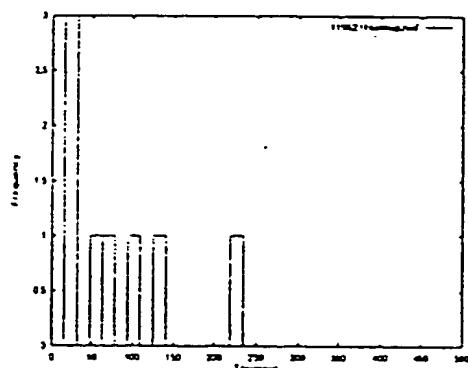
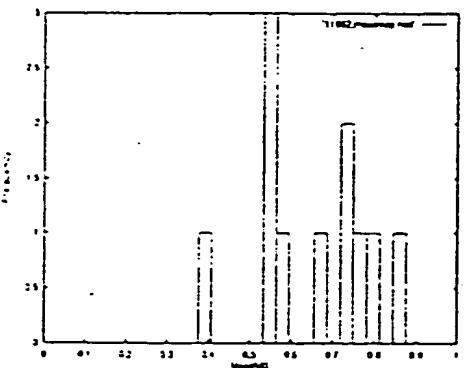
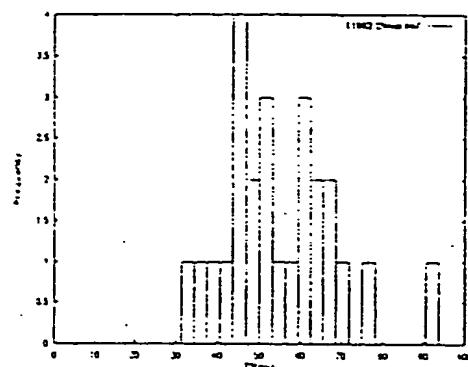
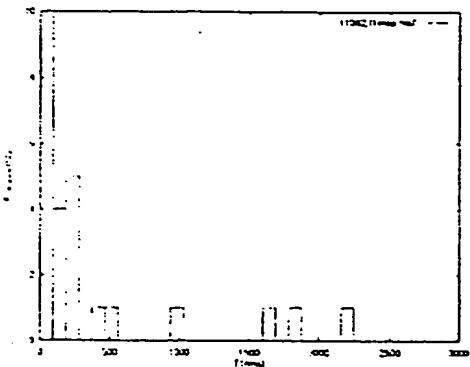
2.1e-06

(s.e.)

0.042

20

6.5e-07



ongg 0004-0009-7

x axis - number of pixels with the same value  
LHS - top T1, middle Msat/M0, bottom M0.

y axis - range of values.

RHS top T2, mid T1sat, bottom ADC

Figure SRL 36 Automated output of histograms for C79 mask 7

T050410-020822860

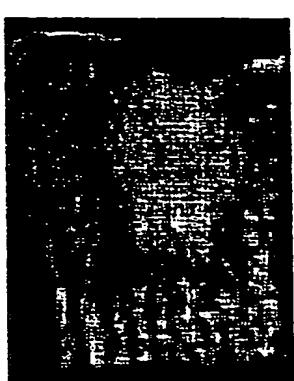
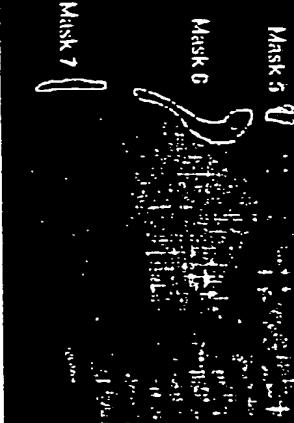
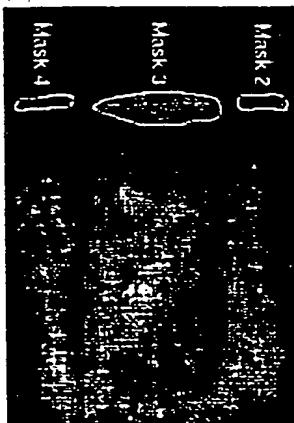
		
C1 Histology 	C1 MRI 	C1 detail of masks 5, 6 and 7 
C2 Histology	C2 MRI not available	C2 detail of masks 2, 3 and 4 
		C1 mask 6 C2 mask 3

Figure SRL 37

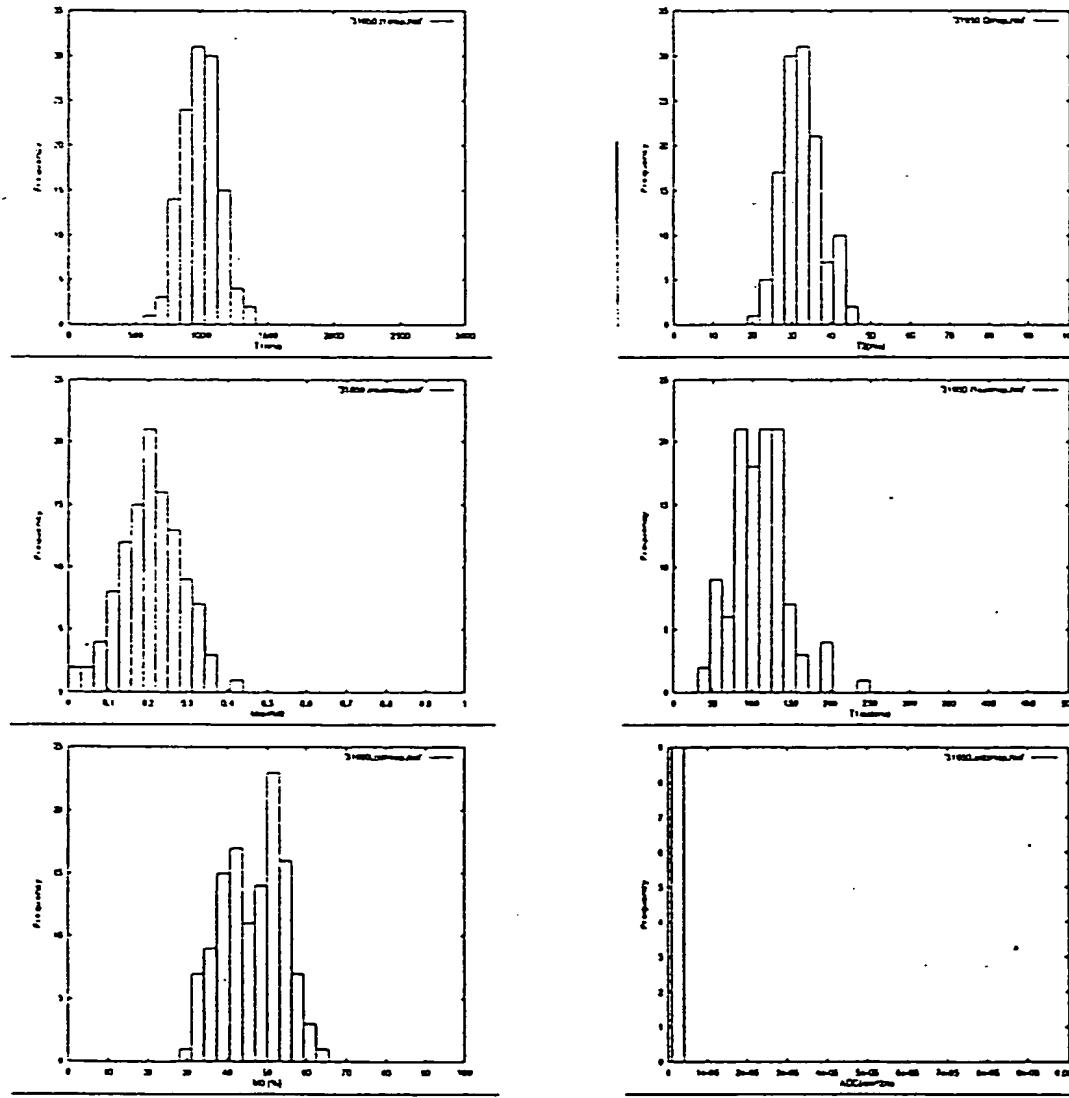
Histology and MRI of spontaneously healed ACD3 defects C1 and C2 at 8 weeks

09828070 . 040501

## C2 mask 3, repair tissue

Parameter	Value	(s.e.)
M0(%)	48.0	0.7
T1(ms)	1028	13
T2(ms)	33.7	0.5

Parameter	Value	(s.e.)
Msat/M0	0.217	0.007
T1sat(ms)	113	3
ADC(cm <sup>2</sup> /s)	1.7e-06	3.2e-07



longg 3804-3809-3

x axis - number of pixels with the same value  
LHS- top T1, middle Msat/M0, bottom M0.

y axis - range of values.

RHS - top T2, mid T1sat, bottom ADC

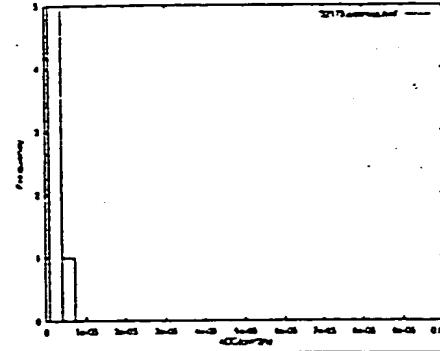
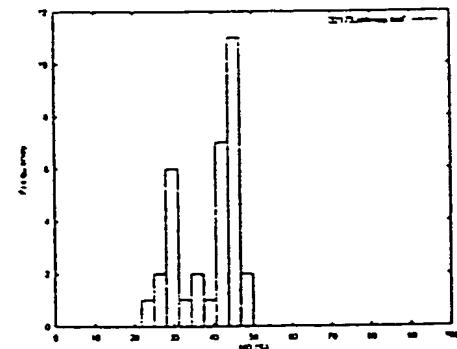
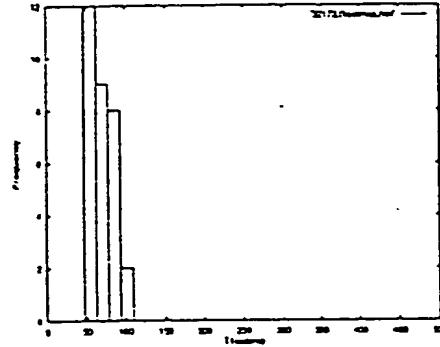
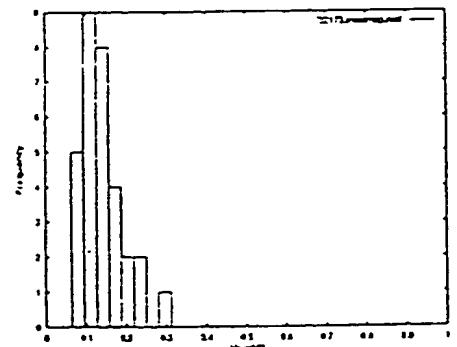
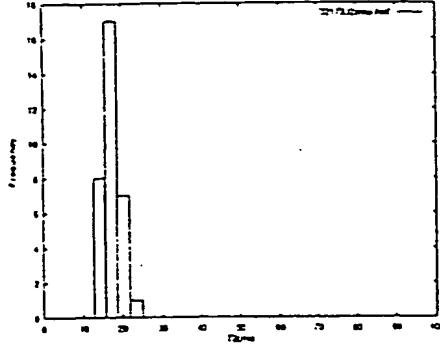
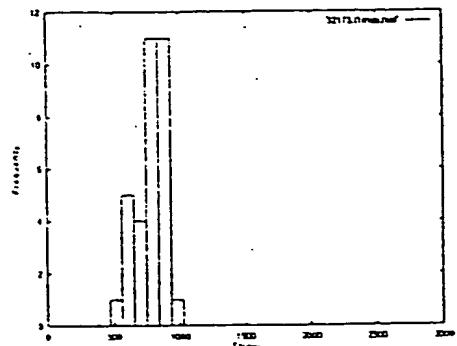
Figure SRL 38 Automated output of histograms for C2 mask 3

00026070 "040507

## C2 mask 4, cartilage adjacent to the defect

Parameter	Value	(s.e.)
M0(%)	40.3	1.3
T1(ms)	816	21
T2(ms)	17.9	0.4

Parameter	Value	(s.e.)
Msat/M0	0.144	0.009
T1sat(ms)	71	3
ADC(cm <sup>2</sup> /s)	1.9e-06	5.9e-07



longg 3804-3809

x axis - number of pixels with the same value  
LHS- top T1, middle Msat/M0, bottom M0.

y axis - range of values.

RHS - top T2, mid T1sat, bottom ADC

Figure SRL 39 Automated output of histograms for C2 mask 4

T050400-02028860

## C1 mask 5, cartilage adjacent to the defect

Parameter Value (s.e.)

M0(%) 48.6 1.5

T1(ms) 830 37

T2(ms) 22.5 1.4

Parameter Value (s.e.)

Msat/M0 0.136 0.012

T1sat(ms) 84 4

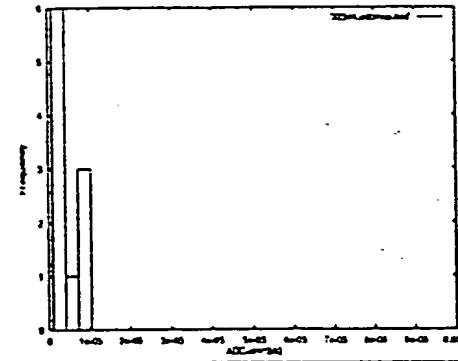
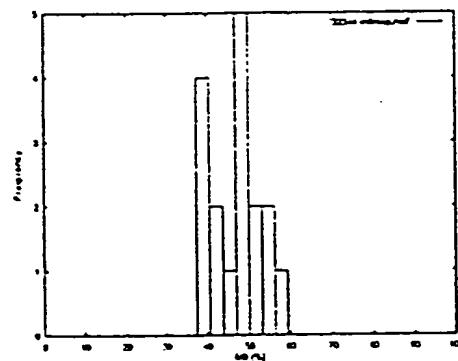
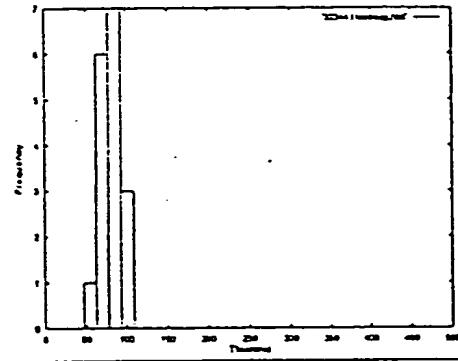
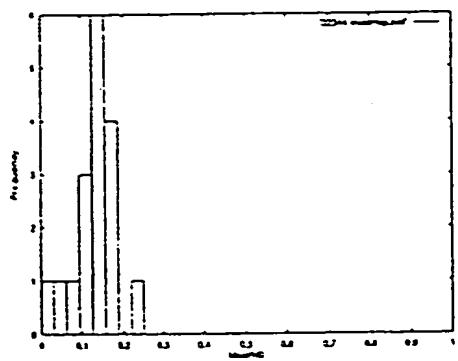
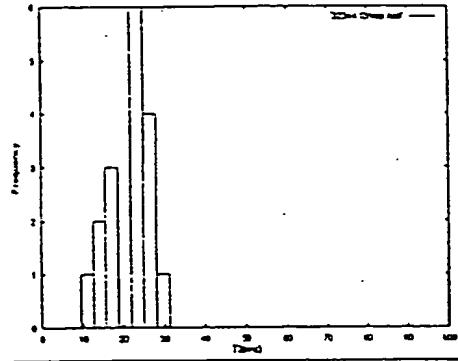
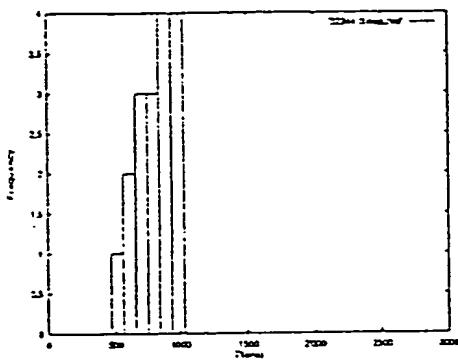
ADC(cm<sup>2</sup>/s) 3.7e-06 7.7e-07

Image ID: 3804-3809-5

x axis - number of pixels with the same value  
LHS- top T1, middle Msat/M0, bottom M0.y axis - range of values.  
RHS - top T2, mid T1sat, bottom ADC

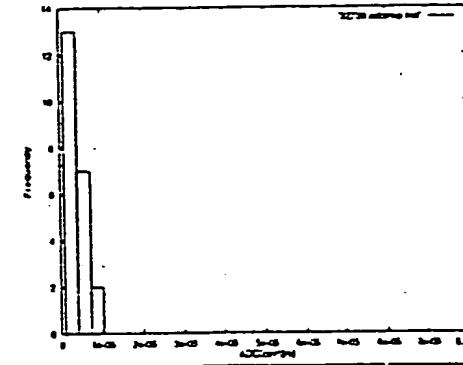
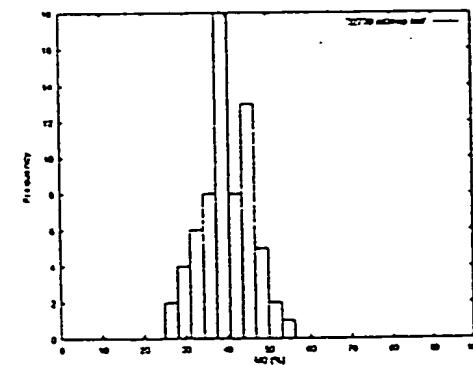
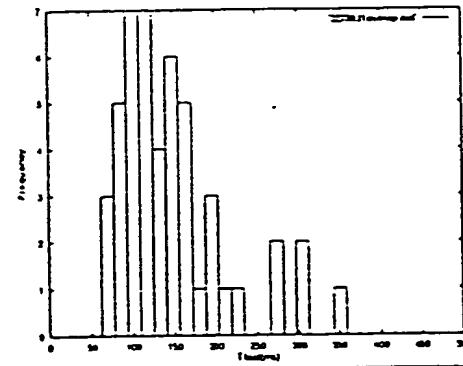
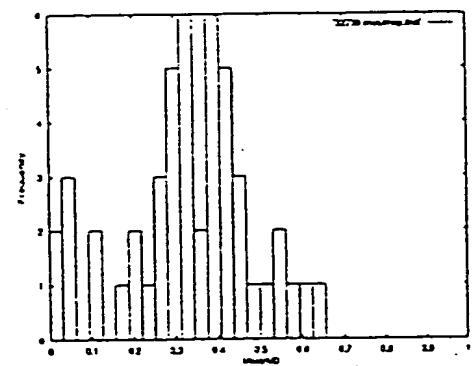
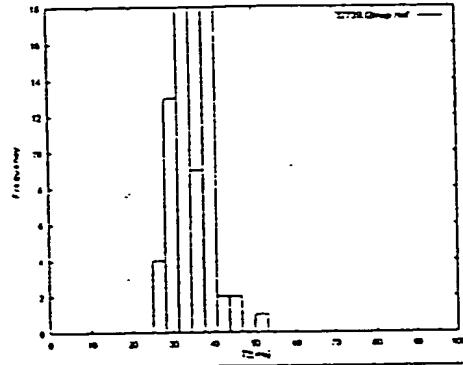
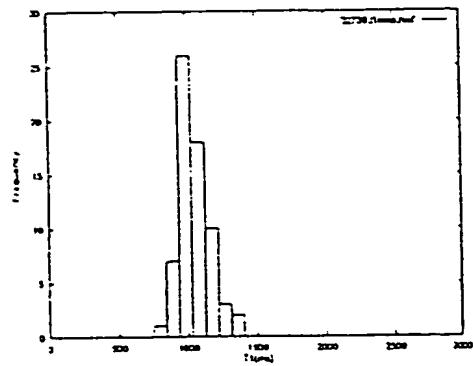
Figure SRL 40 Automated output of histograms for C2 mask 5

Image ID: 3804-3809-5

## C1 mask 6, repair tissue

Parameter	Value	(s.e.)
M0(%)	41.5	0.8
T1(ms)	1082	15
T2(ms)	36.1	0.6

Parameter	Value	(s.e.)
Msat/M0	0.344	0.023
T1sat(ms)	153	10
ADC(cm <sup>2</sup> /s)	3.3e-06	4.5e-07



long 3804-3809

x axis - number of pixels with the same value  
LHS- top T1, middle Msat/M0, bottom M0.

y axis - range of values.

RHS - top T2, mid T1sat, bottom ADC

Figure SRL 41 Automated output of histograms for C1 mask 6

102082860

## C1 mask 7, cartilage adjacent to the defect

## Parameter

M0(%)

T1(ms)

T2(ms)

Value

1.9

22

0.7

(s.e.)

## Parameter

Msat/M0

T1sat(ms)

ADC(cm<sup>2</sup>/s)

Value

0.145

62

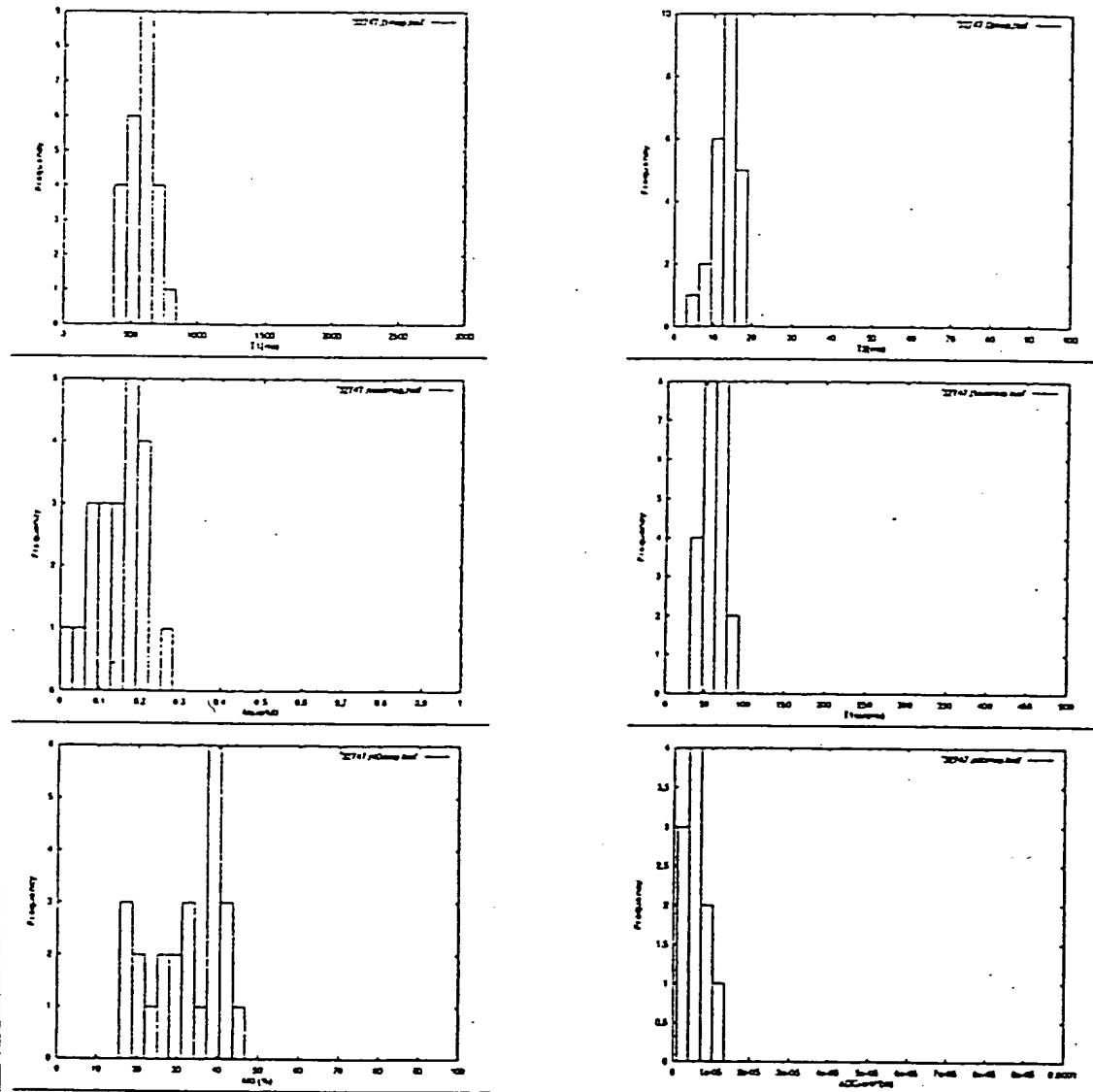
3.9e-06

(s.e.)

0.013

3

9.6e-07



ongg 3804-3809-7

x axis - number of pixels with the same value

y axis - range of values.

LHS- top T1,middleMsat/M0,bottom M0.

RHS - top T2,mid T1sat,bottom ADC

Figure SRL 42 Automated output of histograms for C1 mask 7

09228020 "0405010

C77 (ACD3 + HAED, no synovium) 8 weeks

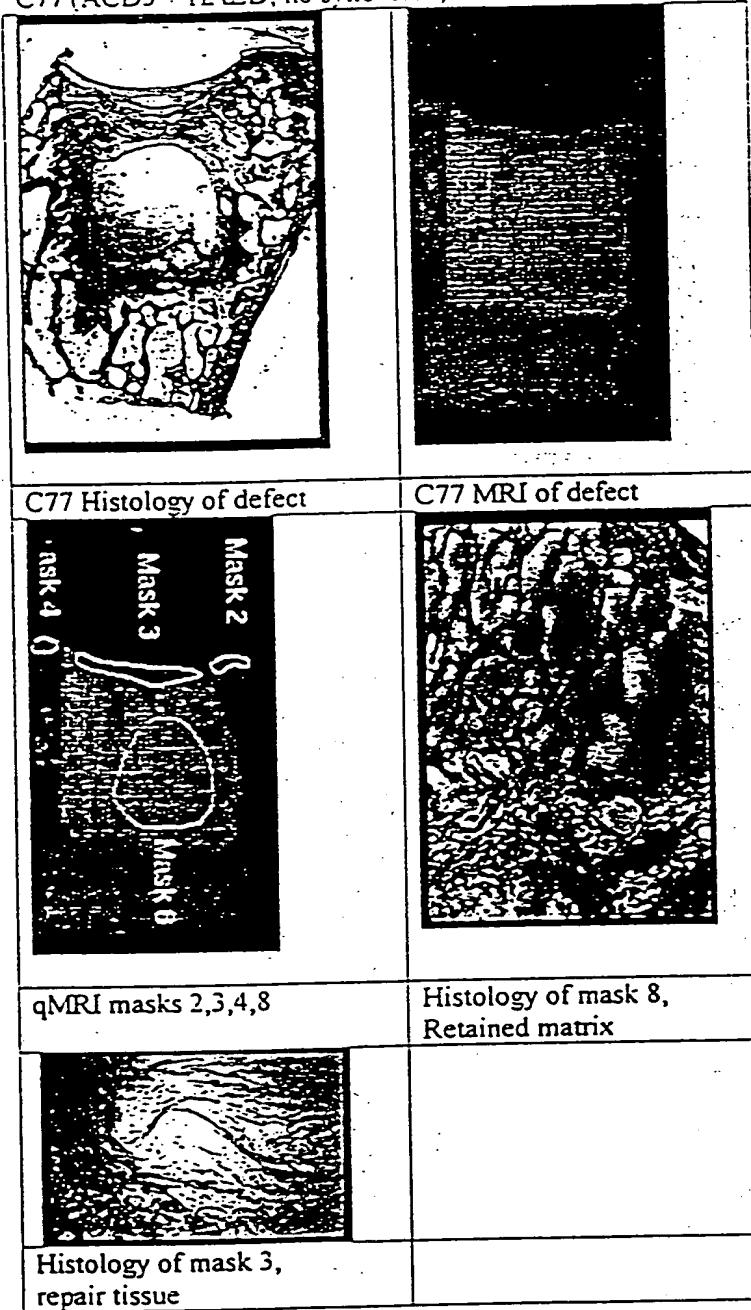


Figure SRL 43

Histology and MRI for C77 (ACD3 + HAED) at 8 weeks

09828070 040501

C77 mask 2

Parameter Value (s.e.)

M0(%) 32.4 1.7

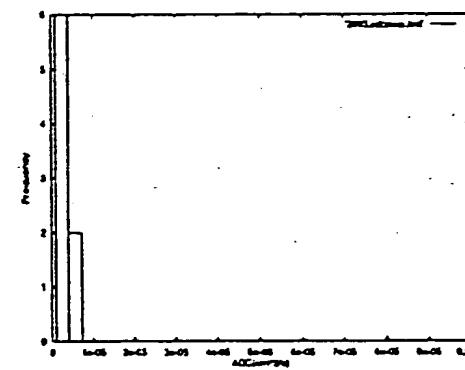
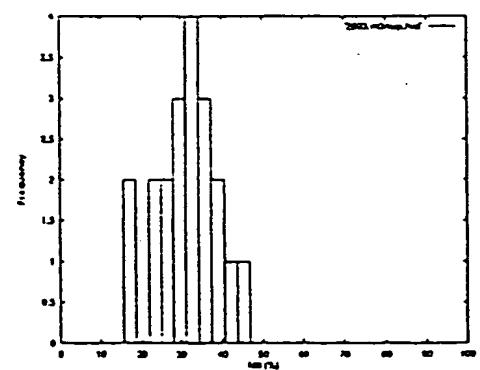
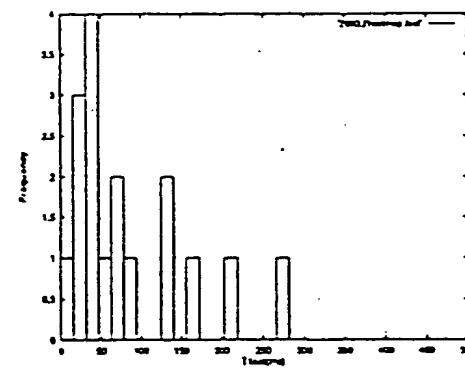
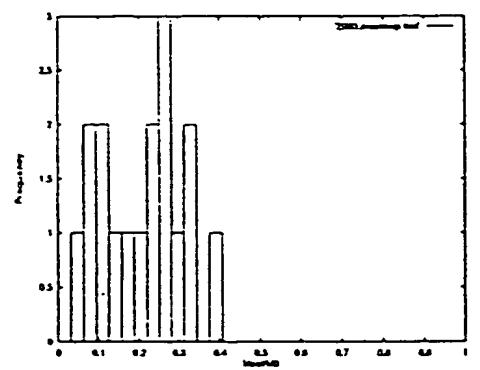
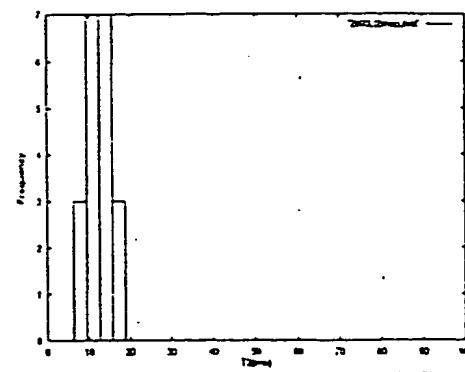
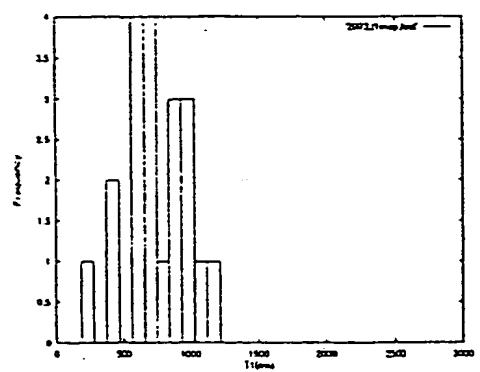
T1(ms) 783 55

T2(ms) 13.2 0.8

Parameter Value (s.e.)

Msat/M0 0.216 0.027

T1sat(ms) 88 19

ADC(cm<sup>2</sup>/s) 3.3e-06 5.6e-07

picc 4413-4418-2

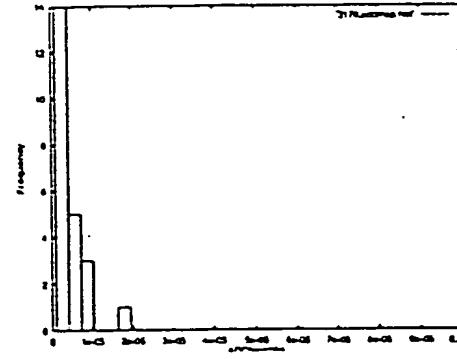
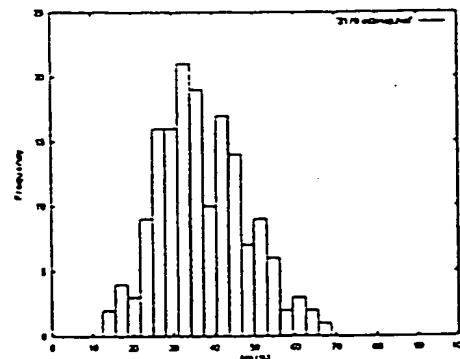
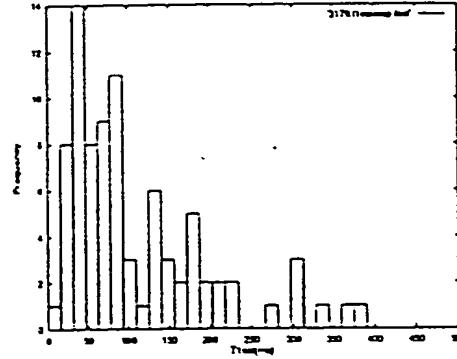
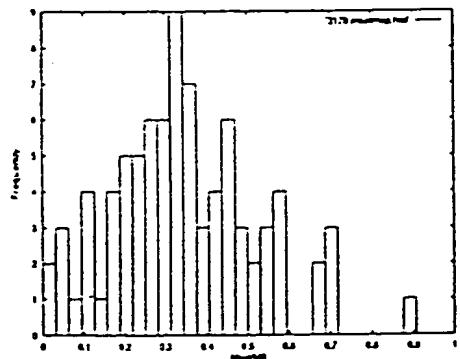
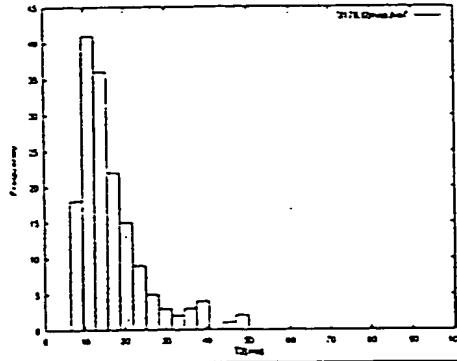
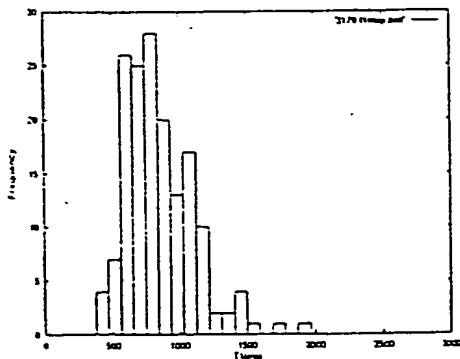
x axis - number of pixels with the same value  
LHS- top T1, middle Msat/M0, bottom M0.y axis - range of values.  
RHS - top T2, mid T1sat, bottom ADC

Figure SRL 44 Automated output of histograms for C77 mask 2

TOP SECRET//NOFORN//EYESAFE

C77 mask 3

Parameter	Value	(s.e.)	Parameter	Value	(s.e.)
M0(%)	38.5	0.9	Msat/M0	0.355	0.020
T1(ms)	884	20	T1sat(ms)	112	10
T2(ms)	17.3	0.7	ADC(cm <sup>2</sup> /s)	2.4e-06	5.1e-07



pic 4413-4418-3

x axis – number of pixels with the same value  
LHS- top T1, middle Msat/M0, bottom M0.

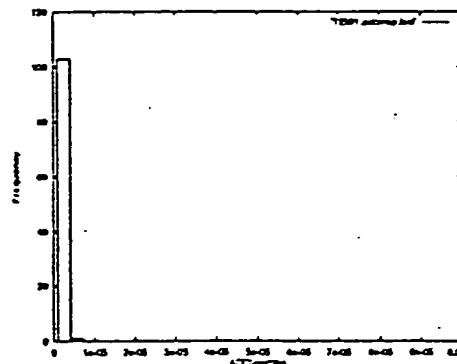
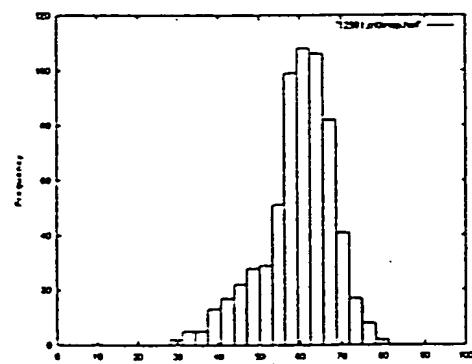
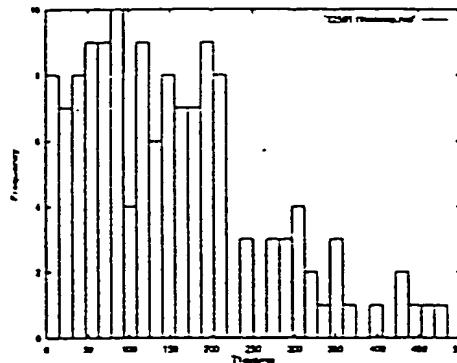
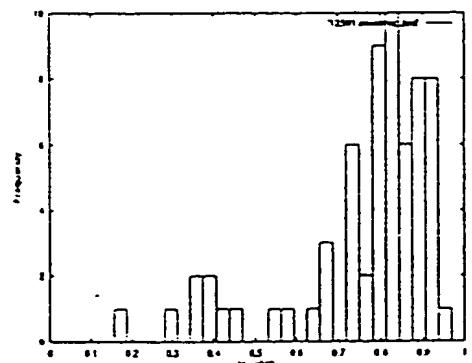
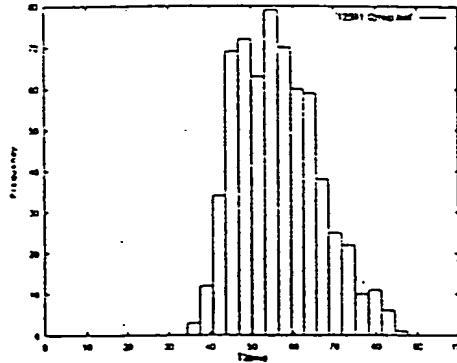
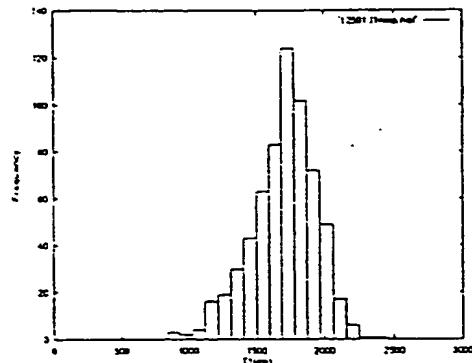
y axis - range of values.

RHS - top T2, mid T1 sat, bottom ADC

Figure SRL 45 Automated output of histograms for C77 mask 3

C77 mask 8

Parameter	Value	(s.e.)	Parameter	Value	(s.e.)
M0(%)	61.4	0.4	Msat/M0	0.785	0.023
T1(ms)	1758	10	T1sat(ms)	222	18
T2(ms)	58.4	0.4	ADC(cm <sup>2</sup> /s)	1.3e-06	6.7e-08



pic 4413-4418-8

x axis - number of pixels with the same value  
LHS- top T1, middle Msat/M0, bottom M0.

y axis - range of values.

RHS - top T2, mid T1 sat, bottom ADC

Figure SRL 46 Automated output of histograms for C77 mask 8

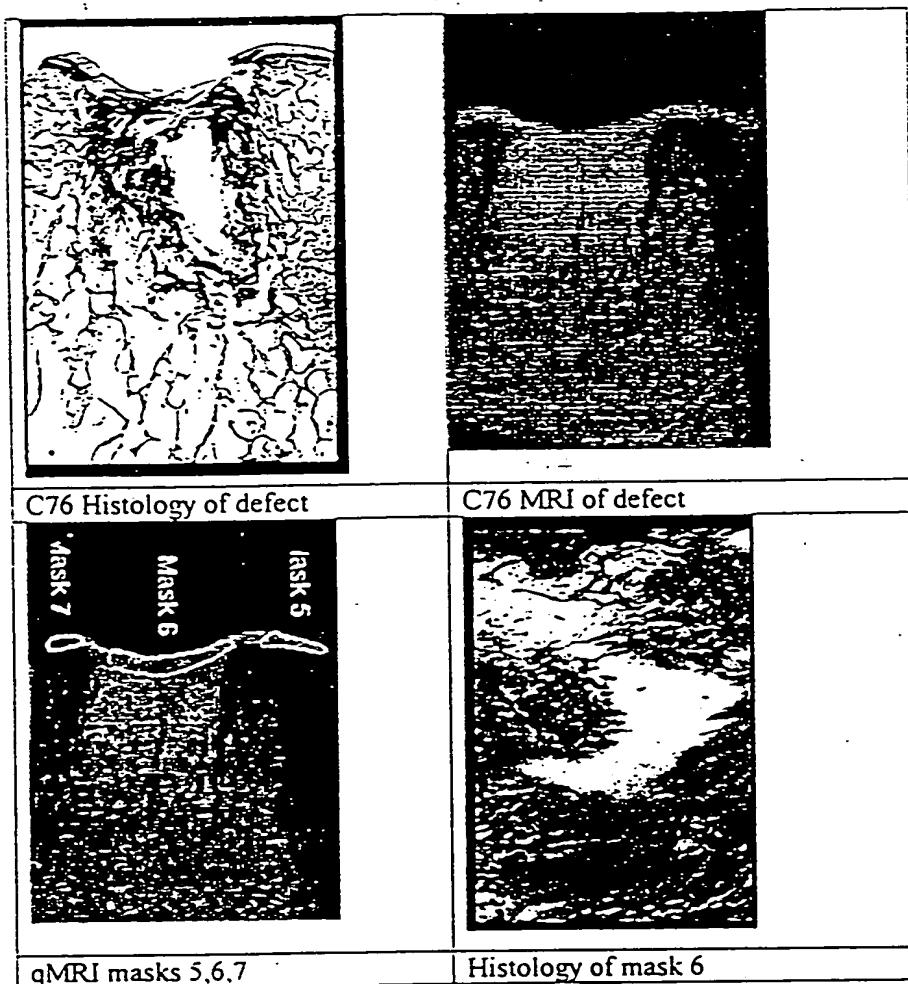


Figure SRL 47  
Histology and MRI for C76 (ACD3 + HAED) at 8 weeks

09828070 040501

## C76 mask 6, retained matrix ÷ repair tissue

## Parameter

M0(%)

62.6

(s.e.)

T1(ms)

1425

26

T2(ms)

77.0

2.5

## Parameter

Msat/M0

0.509

(s.e.)

T1sat(ms)

197

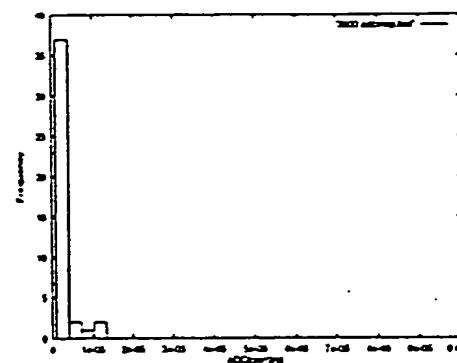
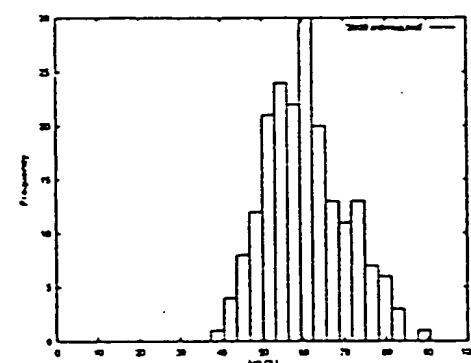
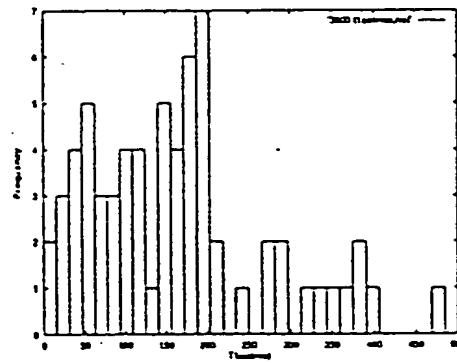
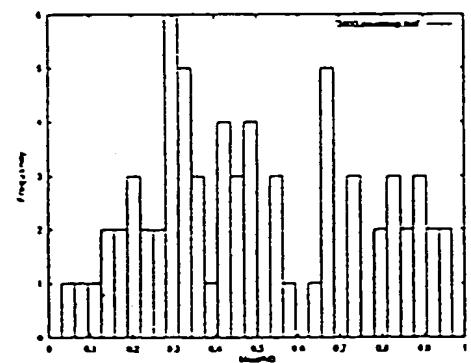
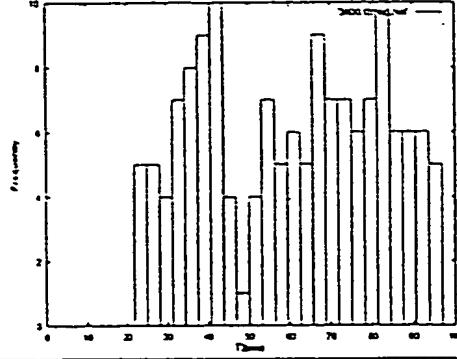
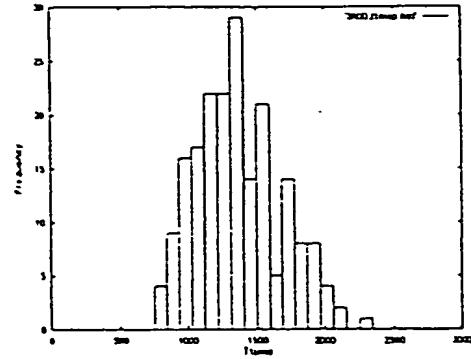
20

ADC(cm<sup>2</sup>/s)

1.8e-06

(s.e.)

2.5e-07



picc 4413-4418-6

x axis - number of pixels with the same value  
LHS- top T1, middle Msat/M0, bottom M0.

y axis - range of values.

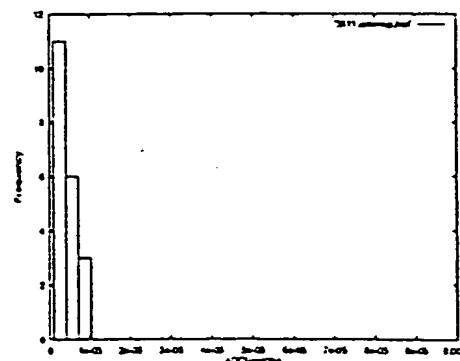
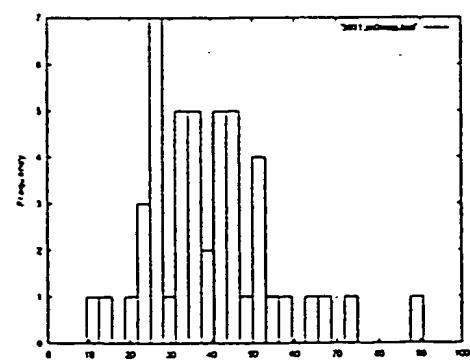
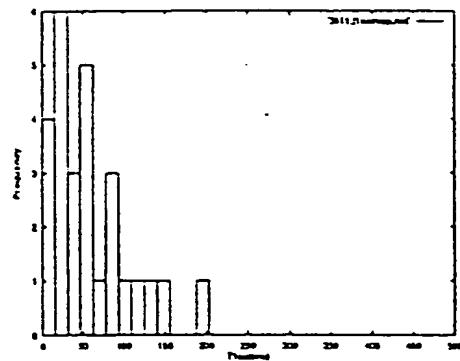
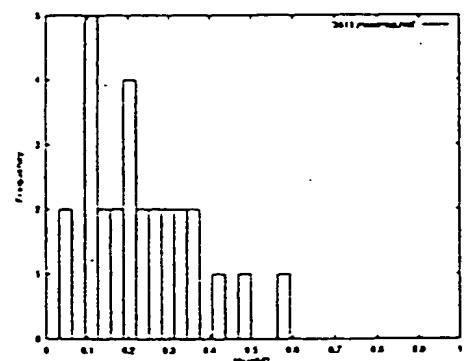
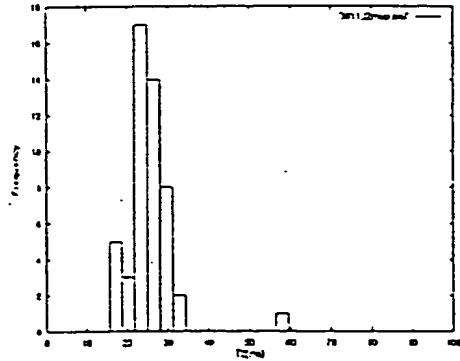
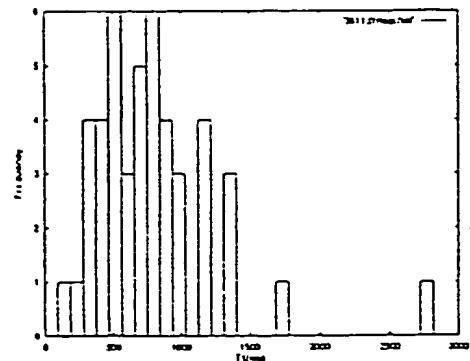
RHS - top T2, mid T1sat, bottom ADC

Figure SRL 48 Automated output of histograms for C76 mask 6

T050400 02020000

C76 mask 5, cartilage adjacent to the defect

Parameter	Value	(s.e.)	Parameter	Value	(s.e.)
M0(%)	40.7	2.3	Msat/M0	0.241	0.025
T1(ms)	864	83	T1sat(ms)	62	10
T2(ms)	26.3	0.9	ADC(cm <sup>2</sup> /s)	3e-06	4.7e-07



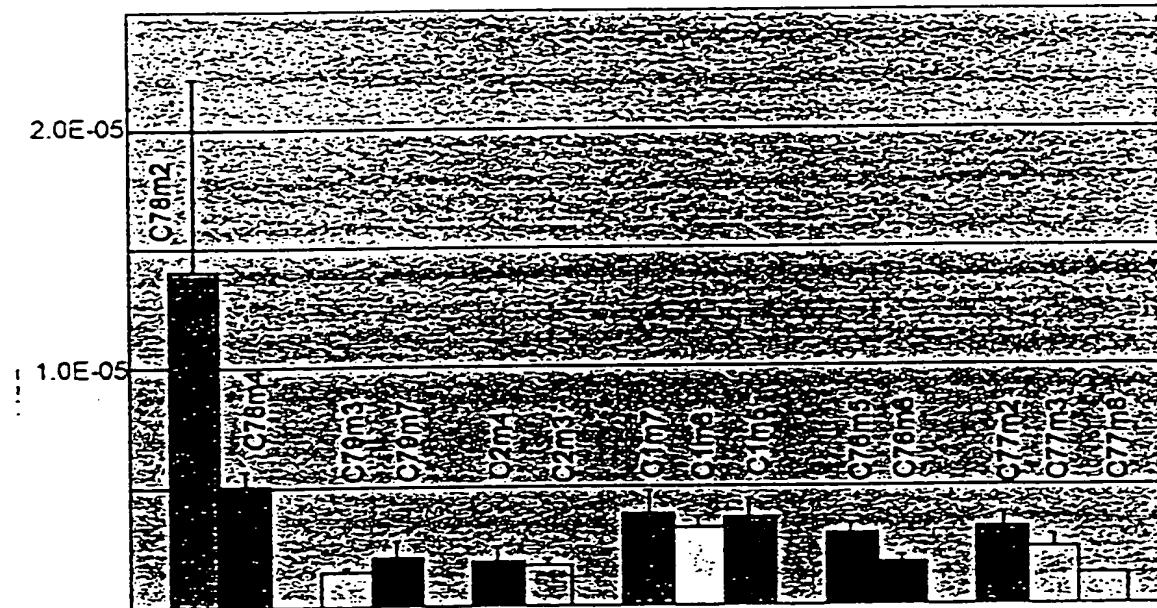
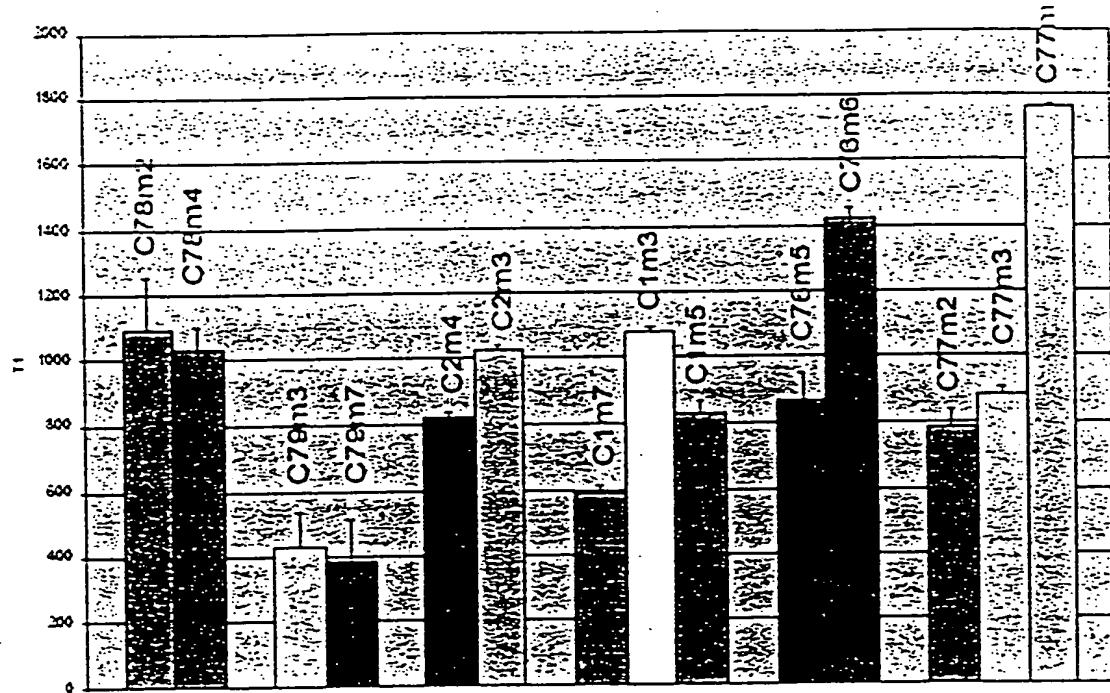
pic 4413-4418-5

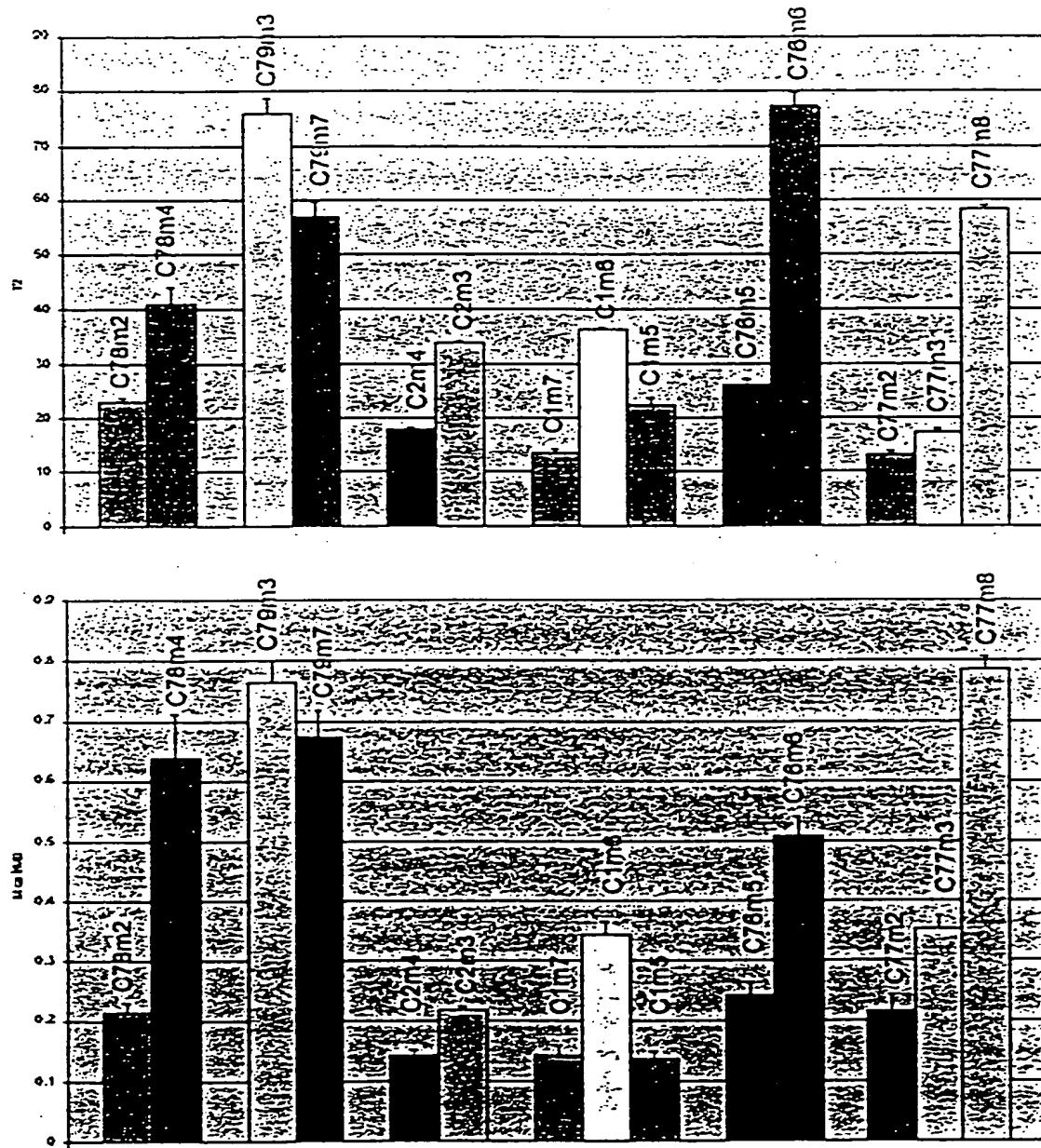
x axis – number of pixels with the same value  
LHS- top T1, middle Msat/M0, bottom M0.

y axis - range of values.

RHS - top T2, mid T1 sat, bottom ADC

Figure SRL 49 Automated output of histograms for C76 mask 5





Summary of  $T_2$  and  $M_{sat}/M_0$  in ACD3 defects

Figure SRL 50b. Summary of mean values of  $T_2$  (above) and  $M_{sat}/M_0$  (below) for repair tissue in ACD3 defects C1, C2, C76, C77, and C79.

7050400 020822360

3D data set C137 slices 1-16

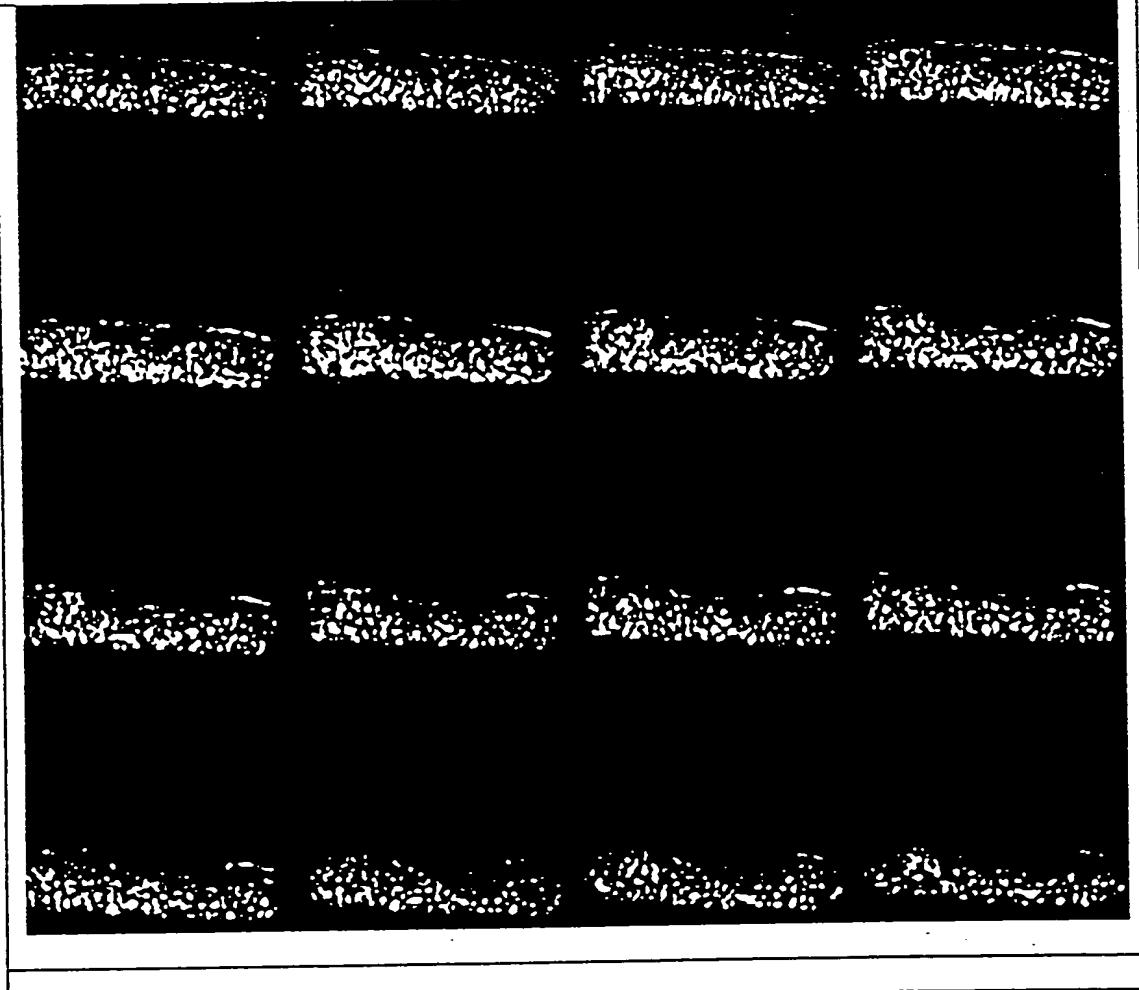


Figure SRL 51a

Adjacent slices (150 $\mu$ m) from a 3D data set showing views from one side of defect C137 (virtual ACD1 + PEG Triacrylate) to the other

3D data set C137 slices 17-32

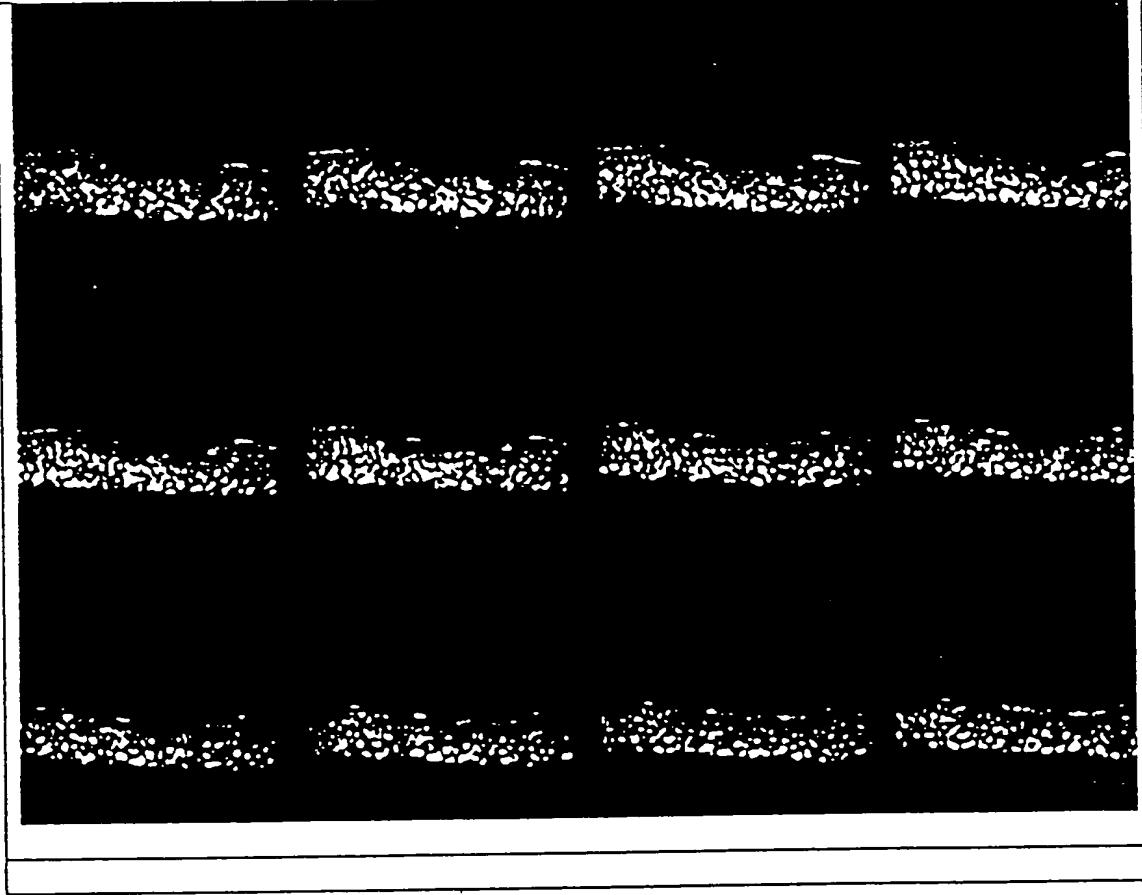


Figure SRL 51b

Adjacent slices (150 $\mu$ m) from a 3D data set showing views from one side of defect C137 (virtual ACD1 + PEG Triacrylate) to the other

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3D data set C137 slices 33-42

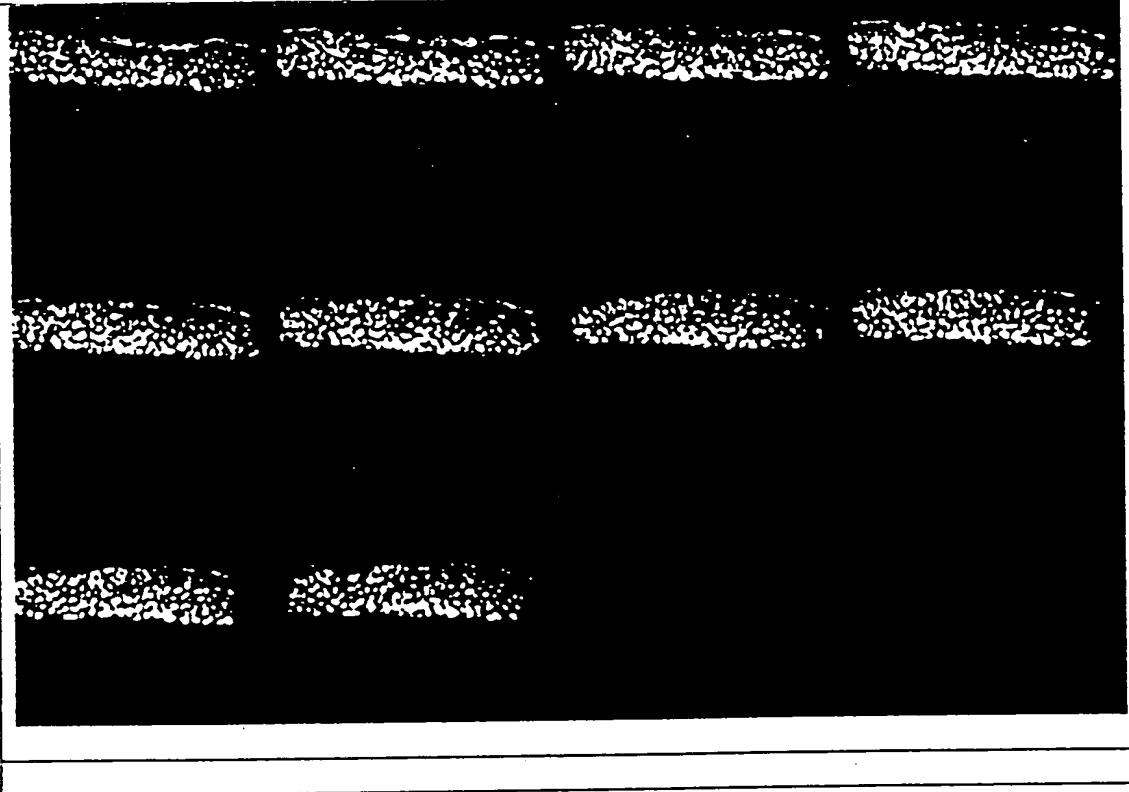


Figure SRL 51c

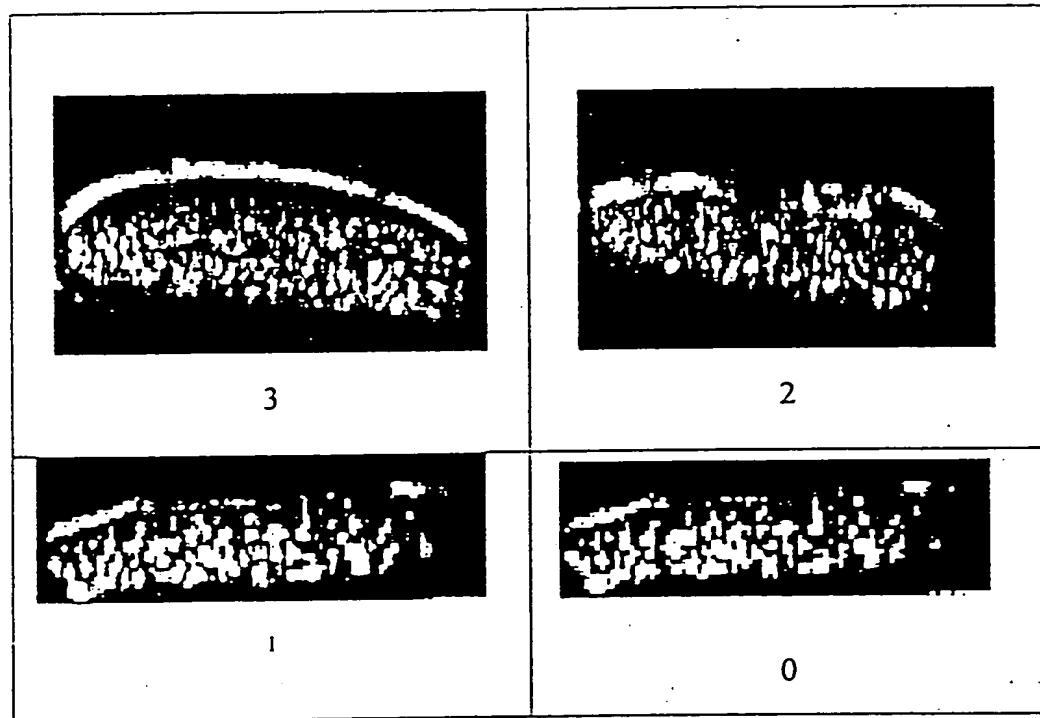
Adjacent slices (150 $\mu$ m) from a 3D data set showing views from one side of defect C137 (virtual ACD1 + PEG Triacrylate) to the other

FO50100-D20082850

## Scoring ACD1 defects from 3-D MRI scans

### Degree of filling of the repair tissue in the defect

- 0 0-20% filling.
- 1 21-40% filling.
- 0 41-70% filling
- 1 70-100% filling.



Sagittal view of defects from a 3D data set

Figure SRL 52

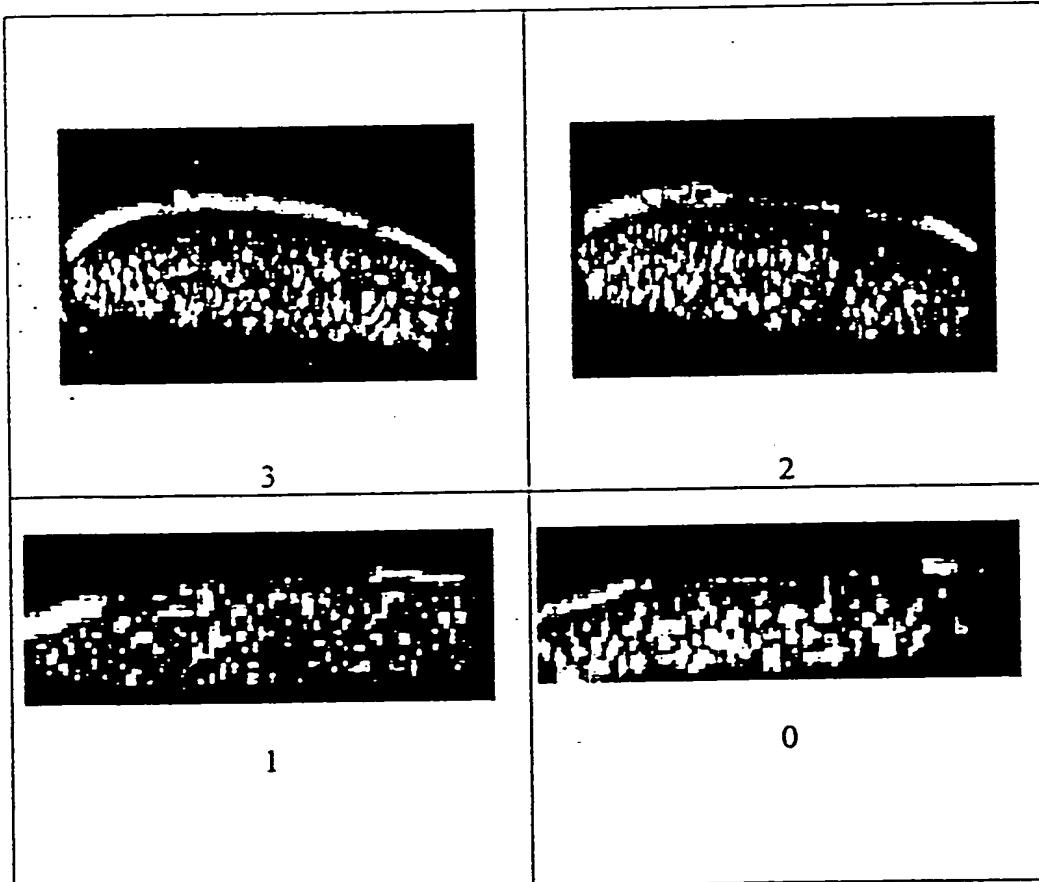
Scoring ACD1 defects from 3-D MRI scans. Degree of filling in the defect

20250410 02082860

## Scoring ACD1 defects from 3-D MRI scans

### Integration of the repair tissue to the surrounding cartilage

- 0 No integration to the sides or bone-cartilage interface.
- 1 Minimal integration (20-40%) to the sides or bone-cartilage interface.
- 2 Moderate integration (41-90%) to the sides or bone-cartilage interface.
- 3 Full integration (91-100%) to the sides and bone-cartilage interface.



Sagittal view of defects from a 3D data set

Figure SRL 53

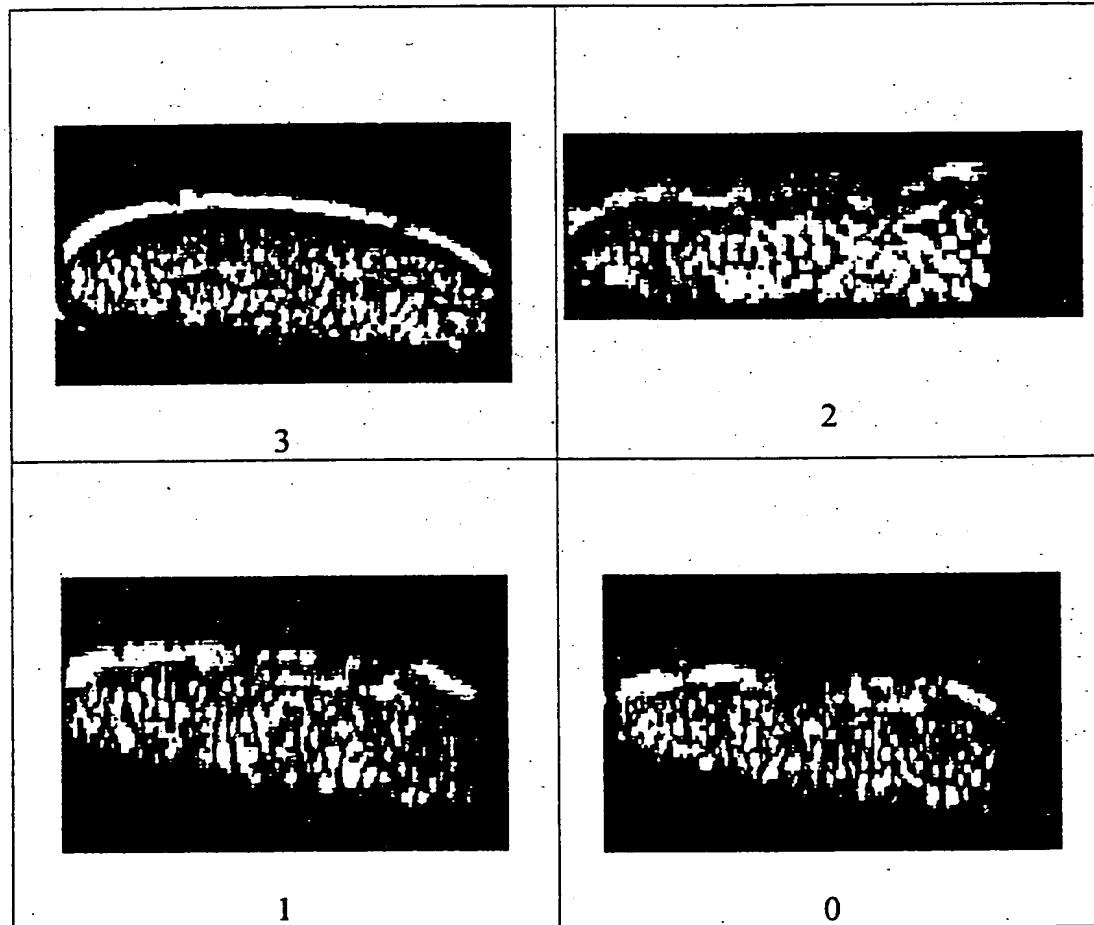
Scoring ACD1 defects from 3-D MRI scans. Integration of repair tissue in the defect

20080408-020822860

## Scoring ACD1 defects from 3-D MRI scans

### Surface smoothness of the repair tissue

- 0 Marked fraying extending to the bone surface in the repair tissue.
- 1 Fraying limited to within the substance of the repair tissue.
- 2 No fraying but the surface appears rough and wavy.
- 3 Smooth surface.



Sagittal view of defects from a 3D data set

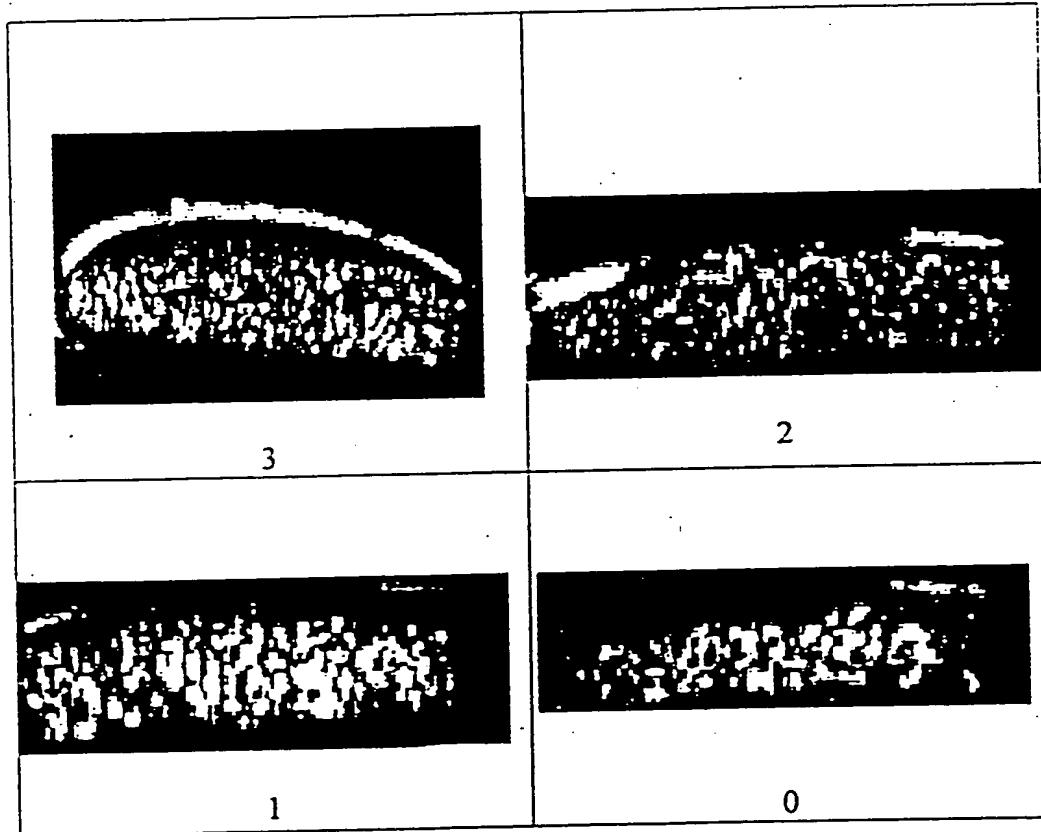
Figure SRL 54

Scoring ACD1 defects from 3-D MRI scans. Smoothness of surface repair

## Scoring ACD1 defects from 3-D MRI scans

### Disruption of subchondral and trabecular bone

- 0 SC bone plate absent with marked loss of trabecular bone.
  - 1 SC bone plate broken with disruption of the trabecular bone pattern.
  - 2 SC bone plate interrupted with no alteration to the trabecular bone pattern.
  - 3 SC bone plate normal with normal trabecular bone underneath.



### Sagittal view of defects from a 3D data set

Figure SRL 55. Scoring ACDI defects from 3-D MRI scans. Disruption of trabecular and subchondral bone

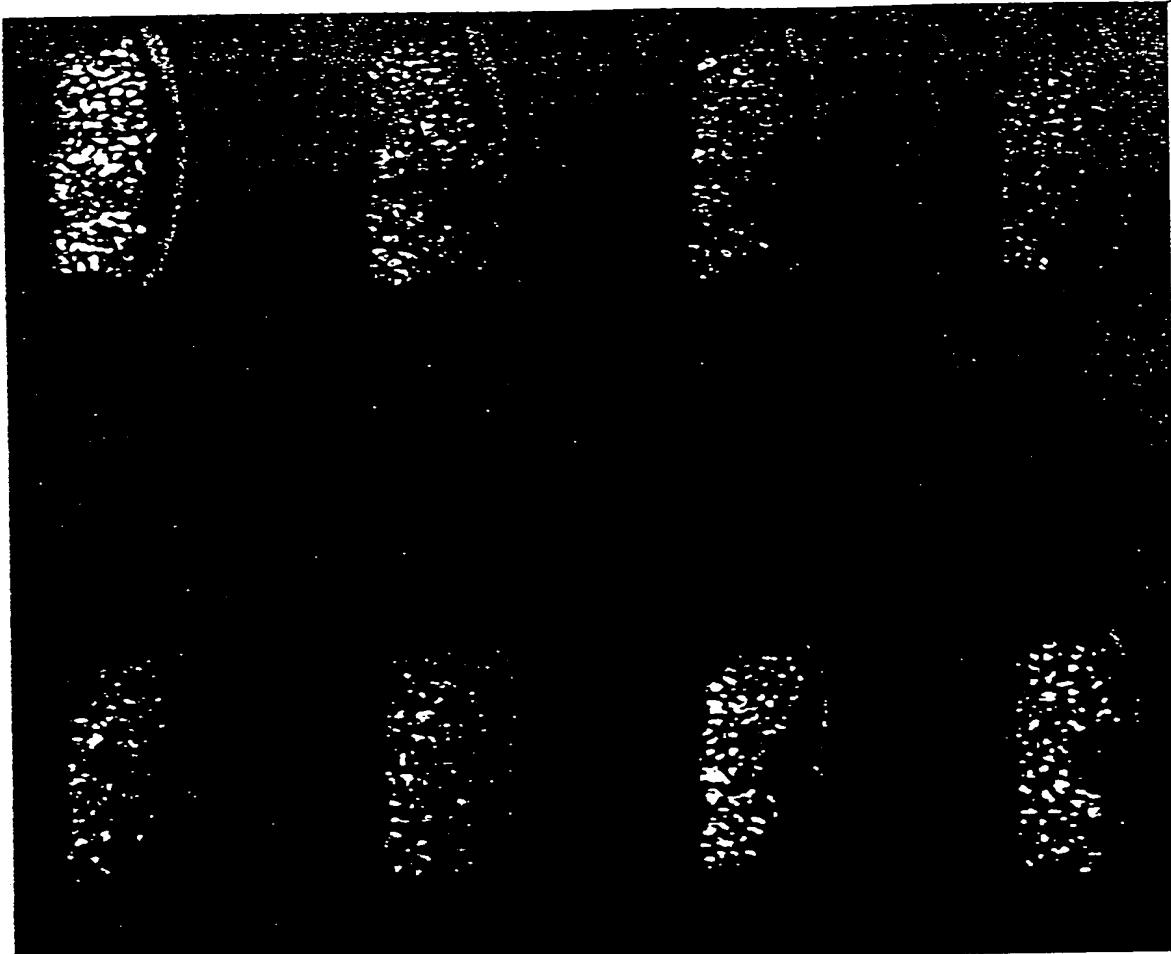
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## Scoring ACD1 defects from 3D MRI scans

G	Specimen	Filling	Integration	Surface	Bone	Signal	Total
G84	C151	3	3	2	3	2	13
G45	C118	0	0	0	3	0	3
G80	C123	3	1	2	2	2	10
+G69	C132	0	0	0	1	0	1
G70	C133	0	0	0	1	0	1
G63	C129	3	2	2	3	2	12
G81	C126	0	0	0	2	0	2
G77	C121	0	0	0	3	0	3
G56	C119	0	0	0	3	1	4
G68	C134	0	0	0	1	0	1
G61	C137	0	0	0	2	1	3
G78L	C139	1	1	1	2	2	7
G78R	C138	0	1	0	2	1	4
G73	C143	2	1	1	3	1	8
G66	C145	0	0	0	2	1	3
G83	C146	0	0	0	3	0	3
K2	C149	1	0	0	3	1	5
G86	C156	1	1	0	3	1	6
G87	C157	2	1	1	2	2	8
G52	C116	0	0	0	3	1	4
G58	C131	2	1	1	3	2	9
G59	C128	3	2	1	3	2	11

Figure SRL 56. Table summarising scores of ACD1 defects from 3-D MRI scans.

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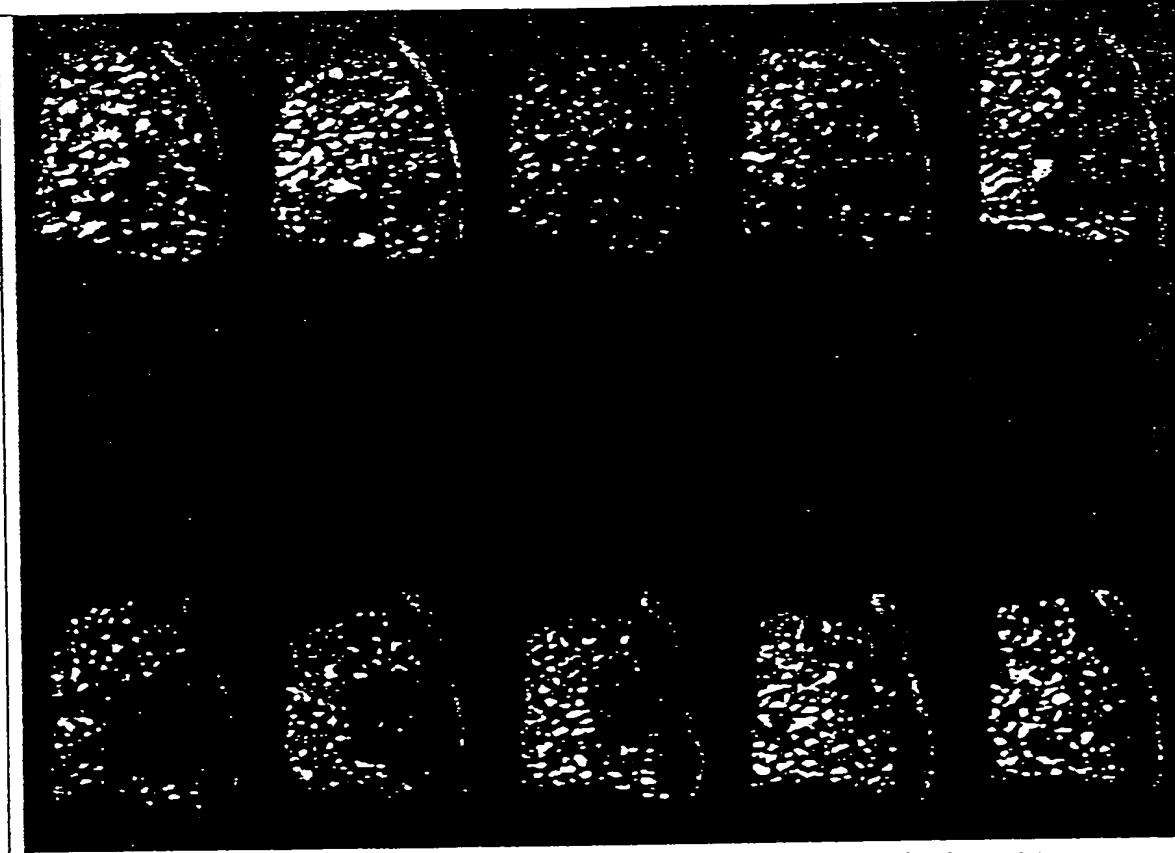


G34 Right medial condyle (C181)

Score : filling -2 , integration-1, surface-2, bone-1, signal-3. Total 9 [max 15]

Figure SRL 57 Slices from a 3D data set of C181 showing views and scoring of an OCT graft in the medial condyle

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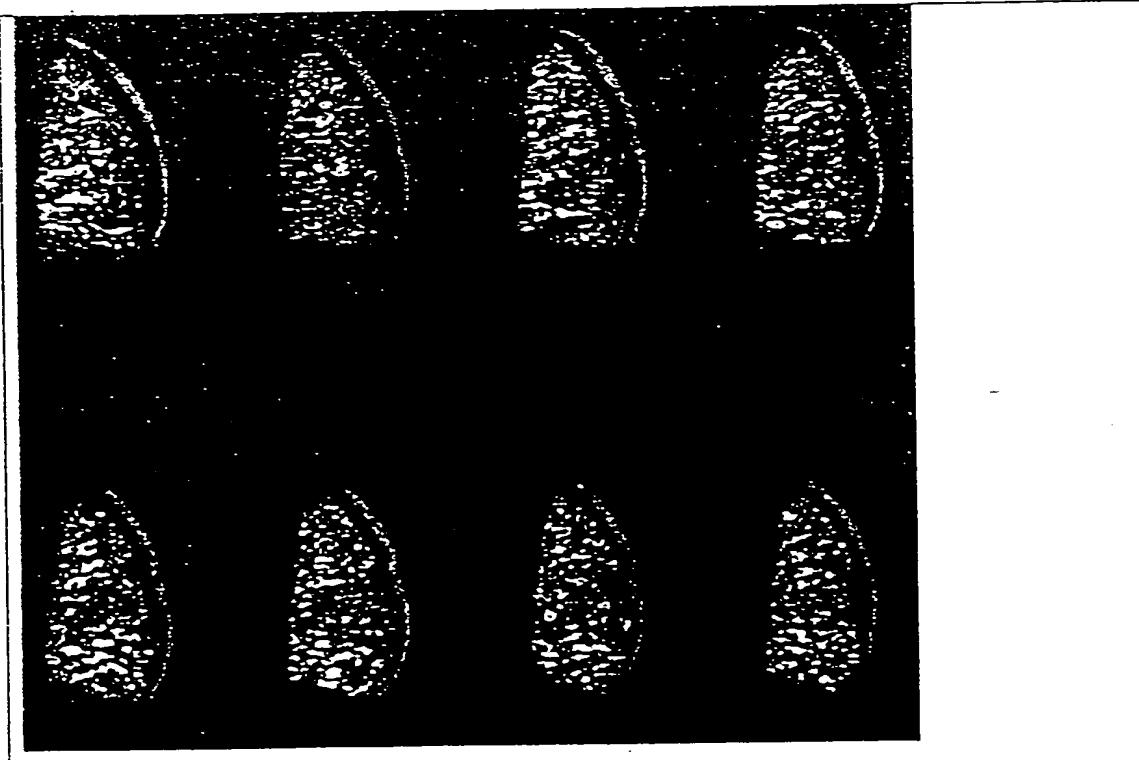


G34 Right lateral condyle (C182)

Score : filling -2 , integration-1, surface-2, bone-1, signal-3. Total 9 [max 15]

10150410-020282860

Figure SRL 58 Slices from a 3D data set of C182 showing views and scoring of an OCT graft in the lateral condyle

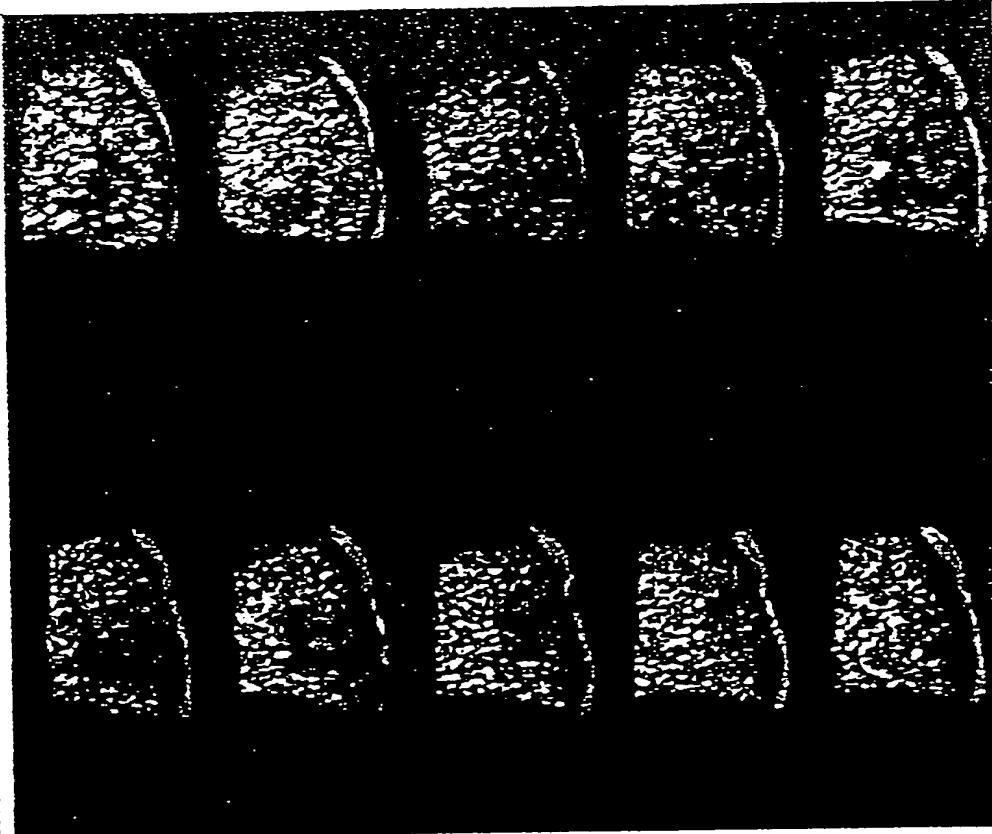


### G33 Right lateral condyle (C174)

Score : filling -3 , integration-3, surface-1, bone-3, signal-3. Total 13 [max 15]

Figure SRL 59 Slices from a 3D data set of C174 showing views and scoring of an OCT graft in the lateral condyle

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### G33 Right medial condyle (C173)

Score : filling -3 , integration-2, surface-2, bone-1, signal-3. Total 11 [max 15]

Figure SRL 60 Slices from a 3D data set of C173 showing views and scoring of an OCT graft in the medial condyle

ప్రాణికాలాలు

## Scoring Osteochondral (OCT) grafts from 3D MRI scans

	Sample	Filling	Integration	Surface	SC Bone	Signal	Total (max 15)
C181	G34 – Rt Knee Medial Condyle	2	1	2	1	3	9
C182	G34 – Rt Knee Lateral Condyle	2	1	2	1	3	9
C173	G33 – Rt Knee Medial Condyle	3	2	2	1	3	11
C174	G33 – Rt Knee Lateral Condyle	3	3	1	3	3	13

Figure SRL 61. Table summarising scores OCT grafts from 3-D MRI scans

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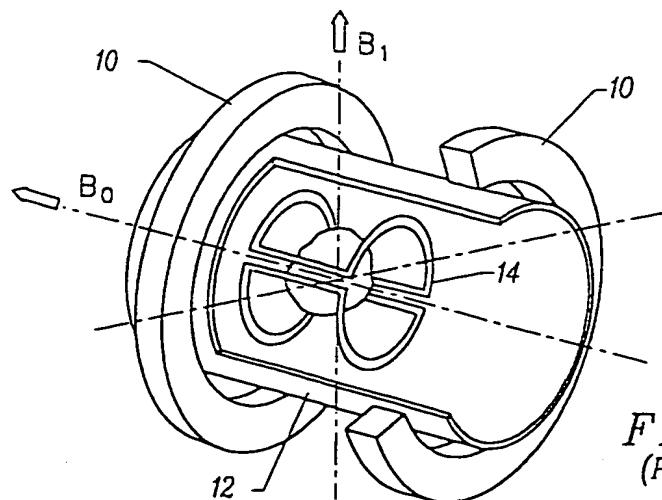
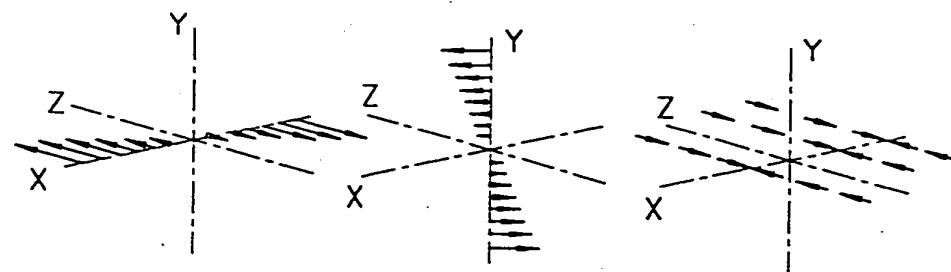
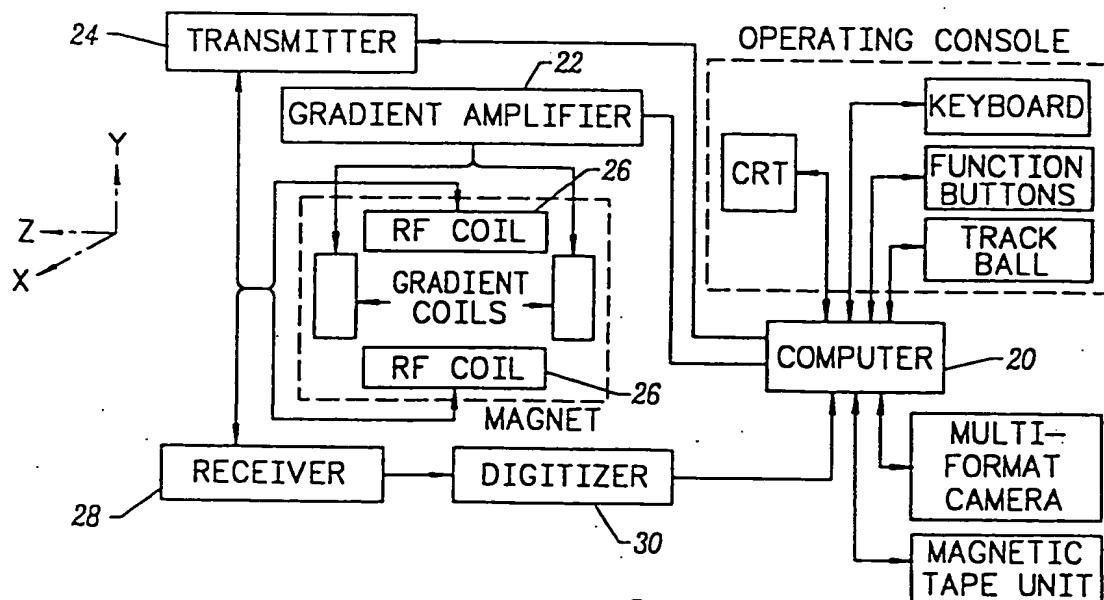
FIG. 61 A  
(PRIOR ART)

FIG. 61 B

FIG. 61 C

FIG. 61 D

FIG. 62  
(PRIOR ART)

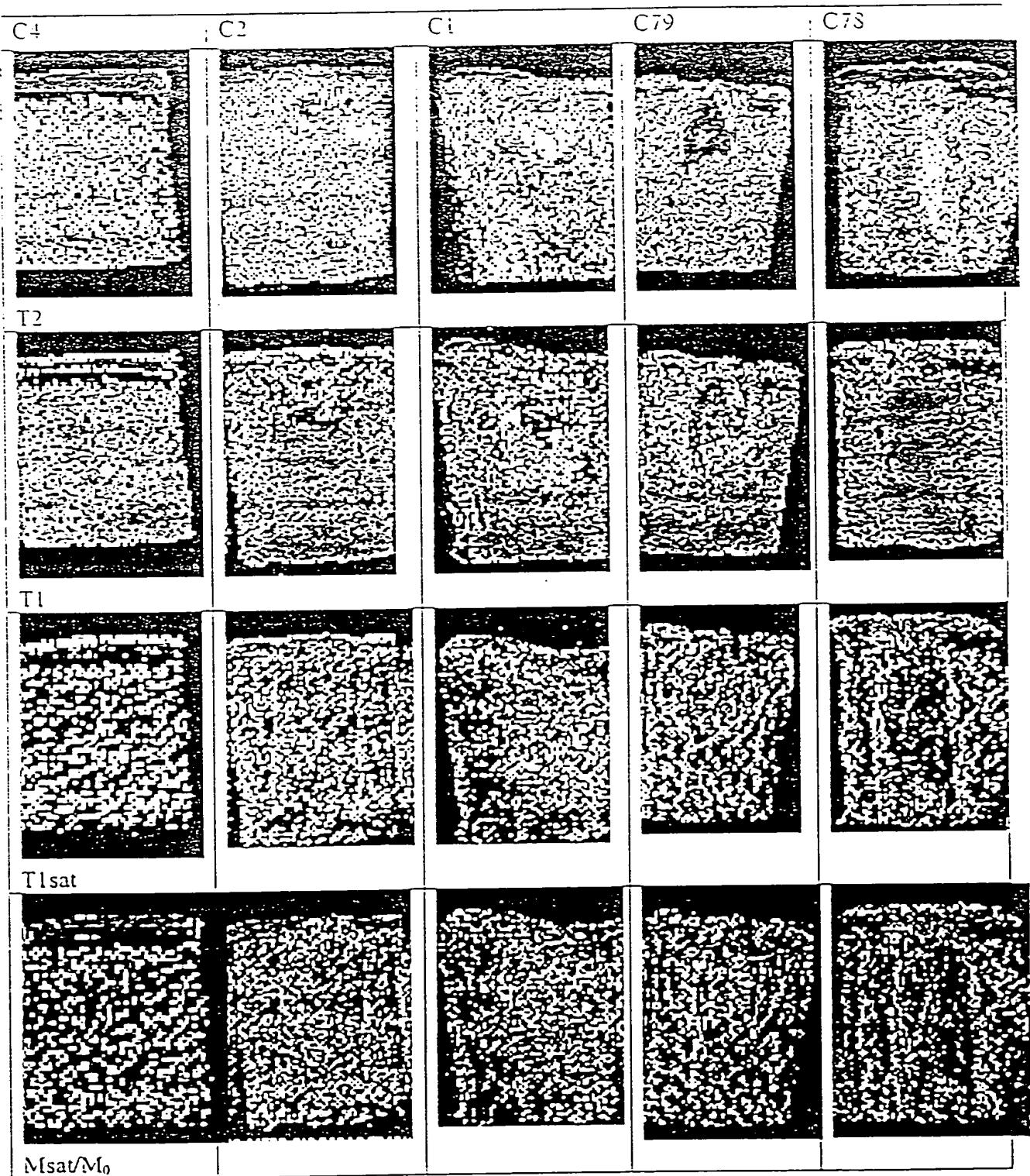
000280270 - 00405001

	C2	C1	C79	C78
T2 contrast (1 <sup>st</sup> echo)				
TR 1500ms TE 6ms				
T2 contrast (last echo)				
TR 1500ms TE 48ms				
T1 contrast				
TR 500ms TE 6ms				
MT contrast				
200ms saturation				

Figure SRL 7

Selected MR scans from qMRI analysis of spontaneously healed ACD3 defects C1, C2, C78, C79 at 8 weeks.

FIG. 64A



Quantitative spatial maps of MR parameters

FIG. 64B

T2 Spatial Maps

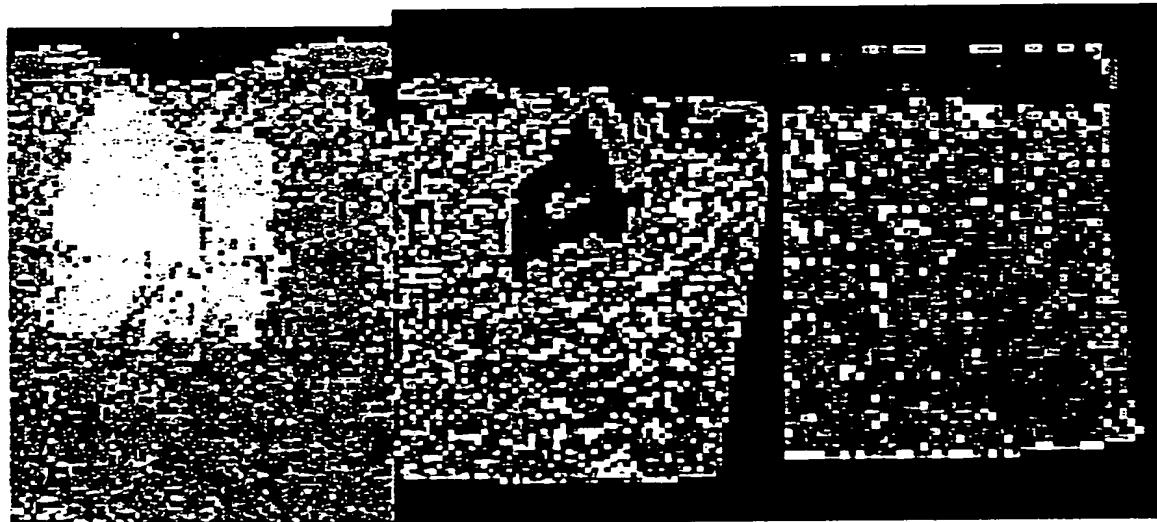
Control cartilage (C4)

ACD3

ACD3

Spontaneous Repair (C1)  
matrix (C77)

Implanted



0932802080-0440561

F16.64C

see Fig. SRL 2  
Figs. SRL 47,48,49

see Figs. SRL 32 & 35

see

T2	T1	T1st	Miss/MQ
1-14 (ms)	1-249 (ms)	1-9 (ms)	0.01-0.09
15-22	250-425	10-29	0.1-0.18
23-30	426-500	30-48	0.19-0.26
31-38	501-775	49-57	0.27-0.34
39-45	776-950	68-86	0.35-0.42
47-54	951-1125	87-105	0.43-0.50
55-62	1126-1300	106-124	0.51-0.58
63-70	1301-1475	125-143	0.59-0.66
71-78	1476-1550	144-152	0.67-0.74
79-86	1651-1825	163-181	0.75-0.82
87-94	1826-2000	182-200	0.83-0.90
95-1000	2001-3000	201-400	0.91-0.99
>1000	>3000	>400	>0.99

Note the colour scales are different for T2 relaxation rates of clinical samples.

Figure 8.8. Range of colours assigned to different qMRI values for spatial maps.

An alternative way of expressing the qMRI data is to colour each pixel according to the range of values shown above. The image can then be displayed as a quantitative spatial map for each MR parameter.

for direct comparison with the histology.

FIG. 65